Robert A Philibert

List of Publications by Year in descending order

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Version: 2024-02-01

188 papers

8,133 citations

47 h-index

47006

80 g-index

188 all docs

188 docs citations

188 times ranked 8285 citing authors

#	Article	IF	CITATIONS
1	The relationship of <i>5HTT</i> (<i>SLC6A4</i>) methylation and genotype on mRNA expression and liability to major depression and alcohol dependence in subjects from the Iowa Adoption Studies. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2008, 147B, 543-549.	1.7	254
2	Methylation Matters: Interaction Between Methylation Density and Serotonin Transporter Genotype Predicts Unresolved Loss or Trauma. Biological Psychiatry, 2010, 68, 405-407.	1.3	242
3	Coordinated changes in AHRR methylation in lymphoblasts and pulmonary macrophages from smokers. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2012, 159B, 141-151.	1.7	230
4	Interplay of genes and early mother–child relationship in the development of selfâ€regulation from toddler to preschool age. Journal of Child Psychology and Psychiatry and Allied Disciplines, 2009, 50, 1331-1338.	5.2	217
5	Methylation at <i>SLC6A4</i> is linked to family history of child abuse: An examination of the Iowa Adoptee sample. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2010, 153B, 710-713.	1.7	209
6	The effect of smoking on DNA methylation of peripheral blood mononuclear cells from African American women. BMC Genomics, 2014, 15, 151.	2.8	193
7	Methylation at 5HTT Mediates the Impact of Child Sex Abuse on Women's Antisocial Behavior: An Examination of the Iowa Adoptee Sample. Psychosomatic Medicine, 2011, 73, 83-87.	2.0	168
8	Prevention Effects Moderate the Association of $5\hat{a}\in HTTLPR$ and Youth Risk Behavior Initiation: Gene $\hat{a}\in f\tilde{A}-\hat{a}\in f$ Environment Hypotheses Tested via a Randomized Prevention Design. Child Development, 2009, 80, 645-661.	3.0	167
9	Changes in DNA methylation at the aryl hydrocarbon receptor repressor may be a new biomarker for smoking. Clinical Epigenetics, 2013, 5, 19.	4.1	167
10	Associations of the serotonin transporter promoter polymorphism with aggressivity, attention deficit, and conduct disorder in an adoptee population. Comprehensive Psychiatry, 2003, 44, 88-101.	3.1	166
11	Serotonin transporter mRNA levels are associated with the methylation of an upstream CpG island. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2007, 144B, 101-105.	1.7	163
12	Social Environment, Genes, and Aggression. American Sociological Review, 2011, 76, 883-912.	5.2	160
13	Economic hardship and biological weathering: The epigenetics of aging in a U.S. sample of black women. Social Science and Medicine, 2016, 150, 192-200.	3.8	156
14	Demethylation of the aryl hydrocarbon receptor repressor as a biomarker for nascent smokers. Epigenetics, 2012, 7, 1331-1338.	2.7	146
15	MAOA methylation is associated with nicotine and alcohol dependence in women. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2008, 147B, 565-570.	1.7	142
16	Gâ€f×â€fE interaction in the organization of attachment: mothers' responsiveness as a moderator of children's genotypes. Journal of Child Psychology and Psychiatry and Allied Disciplines, 2008, 49, 1313-1320.	5.2	129
17	Children's genotypes interact with maternal responsive care in predicting children's competence: Diathesis–stress or differential susceptibility?. Development and Psychopathology, 2011, 23, 605-616.	2.3	128
18	Relationship of serotonin transporter gene polymorphisms and haplotypes to mRNA transcription. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2005, 136B, 58-61.	1.7	113

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19	Cumulative socioeconomic status risk, allostatic load, and adjustment: A prospective latent profile analysis with contextual and genetic protective factors Developmental Psychology, 2013, 49, 913-927.	1.6	112
20	Discrimination, segregation, and chronic inflammation: Testing the weathering explanation for the poor health of Black Americans Developmental Psychology, 2018, 54, 1993-2006.	1.6	112
21	The Impact of Recent Alcohol Use on Genome Wide DNA Methylation Signatures. Frontiers in Genetics, 2012, 3, 54.	2.3	110
22	Epigenetic Silencing of the Human <i>NOS2</i> Gene: Rethinking the Role of Nitric Oxide in Human Macrophage Inflammatory Responses. Journal of Immunology, 2014, 192, 2326-2338.	0.8	107
23	Effects of Genotype and Child Abuse on DNA Methylation and Gene Expression at the Serotonin Transporter. Frontiers in Psychiatry, 2012, 3, 55.	2.6	106
24	Parenting moderates a genetic vulnerability factor in longitudinal increases in youths' substance use Journal of Consulting and Clinical Psychology, 2009, 77, 1-11.	2.0	102
25	Association of the OPRM1 Variant rs1799971 (A118G) with Non-Specific Liability to Substance Dependence in a Collaborative de novo Meta-Analysis of European-Ancestry Cohorts. Behavior Genetics, 2016, 46, 151-169.	2.1	98
26	The effects of social adversity, discrimination, and health risk behaviors on the accelerated aging of African Americans: Further support for the weathering hypothesis. Social Science and Medicine, 2021, 282, 113169.	3.8	98
27	Child maltreatment moderates the association of MAOA with symptoms of depression and antisocial personality disorder Journal of Family Psychology, 2010, 24, 12-20.	1.3	95
28	Dose-Dependent, K+-Stimulated Efflux of Endogenous Taurine from Primary Astrocyte Cultures Is Ca2+-Dependent. Journal of Neurochemistry, 1988, 51, 122-126.	3.9	84
29	Social Adversity, Genetic Variation, Street Code, and Aggression. Youth Violence and Juvenile Justice, 2012, 10, 3-24.	3.0	84
30	Integrated genetic and epigenetic prediction of coronary heart disease in the Framingham Heart Study. PLoS ONE, 2018, 13, e0190549.	2.5	83
31	Differential susceptibility to parenting among African American youths: Testing the DRD4 hypothesis Journal of Family Psychology, 2010, 24, 513-521.	1.3	78
32	The effect of smoking on <i>MAOA</i> promoter methylation in DNA prepared from lymphoblasts and whole blood. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2010, 153B, 619-628.	1.7	76
33	Methylomic Aging as a Window onto the Influence of Lifestyle: Tobacco and Alcohol Use Alter the Rate of Biological Aging. Journal of the American Geriatrics Society, 2015, 63, 2519-2525.	2.6	76
34	A genome-wide search for chromosomal loci linked to mental health wellness in relatives at high risk for bipolar affective disorder among the Old Order Amish. Proceedings of the National Academy of Sciences of the United States of America, 1998, 95, 15531-15536.	7.1	75
35	Replication and meta-analysis of TMEM132D gene variants in panic disorder. Translational Psychiatry, 2012, 2, e156-e156.	4.8	74
36	The impact of stress on the life history strategies of African American adolescents: Cognitions, genetic moderation, and the role of discrimination Developmental Psychology, 2012, 48, 722-739.	1.6	74

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37	DNA methylation age is accelerated in alcohol dependence. Translational Psychiatry, 2018, 8, 182.	4.8	73
38	A pilot examination of the genome-wide DNA methylation signatures of subjects entering and exiting short-term alcohol dependence treatment programs. Epigenetics, 2014, 9, 1212-1219.	2.7	72
39	Current and Future Prospects for Epigenetic Biomarkers of Substance Use Disorders. Genes, 2015, 6, 991-1022.	2.4	70
40	Behavioral genetics in antisocial spectrum disorders and psychopathy: A review of the recent literature. Behavioral Sciences and the Law, 2010, 28, 148-173.	0.8	67
41	The structure and expression of the human neuroligin-3 gene. Gene, 2000, 246, 303-310.	2.2	63
42	Reversion of AHRR Demethylation Is a Quantitative Biomarker of Smoking Cessation. Frontiers in Psychiatry, 2016, 7, 55.	2.6	58
43	Lower TSH and higher T4 levels are associated with current depressive syndrome in young adults. Acta Psychiatrica Scandinavica, 2006, 114, 132-139.	4.5	55
44	Differential sensitivity to prevention programming: A dopaminergic polymorphism-enhanced prevention effect on protective parenting and adolescent substance use Health Psychology, 2014, 33, 182-191.	1.6	55
45	AHRR methylation predicts smoking status and smoking intensity in both saliva and blood DNA. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2020, 183, 51-60.	1.7	55
46	Harsh parenting and adolescent health: A longitudinal analysis with genetic moderation Health Psychology, 2014, 33, 401-409.	1.6	54
47	Perceived discrimination, serotonin transporter linked polymorphic region status, and the development of conduct problems. Development and Psychopathology, 2011, 23, 617-627.	2.3	53
48	A quantitative epigenetic approach for the assessment of cigarette consumption. Frontiers in Psychology, 2015, 6, 656.	2.1	53
49	Impact of child sex abuse on adult psychopathology: A genetically and epigenetically informed investigation Journal of Family Psychology, 2013, 27, 3-11.	1.3	52
50	Parenting, Socioeconomic Status Risk, and Later Young Adult Health: Exploration of Opposing Indirect Effects via DNA Methylation. Child Development, 2016, 87, 111-121.	3.0	50
51	Prevention Effects Ameliorate the Prospective Association Between Nonsupportive Parenting and Diminished Telomere Length. Prevention Science, 2015, 16, 171-180.	2.6	48
52	Comparison of the genotyping results using DNA obtained from blood and saliva. Psychiatric Genetics, 2008, 18, 275-281.	1.1	47
53	Neighborhood crime and depressive symptoms among African American women: Genetic moderation and epigenetic mediation of effects. Social Science and Medicine, 2015, 146, 120-128.	3.8	47
54	Association between the serotonin transporter promoter polymorphism (5-HTTLPR) and adult unresolved attachment Developmental Psychology, 2009, 45, 64-76.	1.6	44

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55	Methylation of the oxytocin receptor gene mediates the effect of adversity on negative schemas and depression. Development and Psychopathology, 2017, 29, 725-736.	2.3	44
56	Predictors of suicidal ideation, suicide attempts, and self-harm without lethal intent in a community corrections sample. Journal of Criminal Justice, 2011, 39, 238-245.	2.3	43
57	Coordinated DNA methylation and gene expression changes in smoker alveolar macrophages: specific effects on VEGF receptor 1 expression. Journal of Leukocyte Biology, 2012, 92, 621-631.	3.3	43
58	Olanzapine Usage Associated With Neuroleptic Malignant Syndrome. Psychosomatics, 2001, 42, 528-529.	2.5	42
59	Dose Response and Prediction Characteristics of a Methylation Sensitive Digital PCR Assay for Cigarette Consumption in Adults. Frontiers in Genetics, 2018, 9, 137.	2.3	42
60	Association of an X-chromosome dodecamer insertional variant allele with mental retardation. Molecular Psychiatry, 1998, 3, 303-309.	7.9	41
61	Is serotonin transporter genotype associated with epigenetic susceptibility or vulnerability? Examination of the impact of socioeconomic status risk on African American youth. Development and Psychopathology, 2014, 26, 289-304.	2.3	41
62	Accuracy and utility of an epigenetic biomarker for smoking in populations with varying rates of false selfâ€report. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2017, 174, 641-650.	1.7	41
63	Gene environment interactions with a novel variable Monoamine Oxidase A transcriptional enhancer are associated with antisocial personality disorder. Biological Psychology, 2011, 87, 366-371.	2.2	40
64	Sharing the Burden of the Transition to Adulthood: African American Young Adults' Transition Challenges and Their Mothers' Health Risk. American Sociological Review, 2018, 83, 143-172.	5.2	40
65	Role of GABRA2 on risk for alcohol, nicotine, and cannabis dependence in the Iowa Adoption Studies. Psychiatric Genetics, 2009, 19, 91-98.	1.1	39
66	Supportive family environments, genes that confer sensitivity, and allostatic load among rural african american emerging adults: A prospective analysis Journal of Family Psychology, 2013, 27, 22-29.	1.3	39
67	Neighborhood Disadvantage and Biological Aging: Using Marginal Structural Models to Assess the Link Between Neighborhood Census Variables and Epigenetic Aging. Journals of Gerontology - Series B Psychological Sciences and Social Sciences, 2019, 74, e50-e59.	3.9	39
68	Life stress, the dopamine receptor gene, and emerging adult drug use trajectories: A longitudinal, multilevel, mediated moderation analysis. Development and Psychopathology, 2012, 24, 941-951.	2.3	38
69	A Triplet Repeat on 17q Accounts for Most Expansions Detected by the Repeat-Expansion–Detection Technique. American Journal of Human Genetics, 1998, 62, 1548-1551.	6.2	36
70	The relationship of deiodinase 1 genotype and thyroid function to lifetime history of major depression in three independent populations. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2011, 156, 593-599.	1.7	36
71	Methylation array data can simultaneously identify individuals and convey protected health information: an unrecognized ethical concern. Clinical Epigenetics, 2014, 6, 28.	4.1	35
72	Genetically contextual effects of smoking on genome wide DNA methylation. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2017, 174, 595-607.	1.7	34

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73	Population-based association analyses of the HOPA12bp polymorphism for schizophrenia and hypothyroidism. American Journal of Medical Genetics Part A, 2001, 105, 130-134.	2.4	33
74	Participation in a Family-Centered Prevention Program Decreases Genetic Risk for Adolescents' Risky Behaviors. Pediatrics, 2009, 124, 911-917.	2.1	33
75	Alcohol and tobacco consumption alter hypothalamic pituitary adrenal axis DNA methylation. Psychoneuroendocrinology, 2016, 66, 176-184.	2.7	33
76	HPLC analysis of putative amino acid neurotransmitters released from primary cerebellar cultures. Journal of Neuroscience Methods, 1987, 22, 173-179.	2.5	32
77	Investigation of a candidate gene for schizophrenia on Xq13 previously associated with mental retardation and hypothyroidism. American Journal of Medical Genetics Part A, 2000, 96, 398-403.	2.4	31
78	Differential impact of cumulative SES risk on methylation of protein–protein interaction pathways as a function of SLC6A4 genetic variation in African American young adults. Biological Psychology, 2014, 96, 28-34.	2.2	31
79	An index of the ratio of inflammatory to antiviral cell types mediates the effects of social adversity and age on chronic illness. Social Science and Medicine, 2017, 185, 158-165.	3.8	31
80	K+-evoked taurine efflux from cerebellar astrocytes: On the roles of Ca2+ and Na+. Neurochemical Research, 1989, 14, 43-48.	3.3	30
81	Genetic moderation of contextual effects on negative arousal and parenting in African-American parents Journal of Family Psychology, 2012, 26, 46-55.	1.3	30
82	Blood-Based Biomarkers for Predicting the Risk for Five-Year Incident Coronary Heart Disease in the Framingham Heart Study via Machine Learning. Genes, 2018, 9, 641.	2.4	29
83	Role of MED12 in transcription and human behavior. Pharmacogenomics, 2007, 8, 909-916.	1.3	28
84	Childhood/Adolescent stressors and allostatic load in adulthood: Support for a calibration model. Social Science and Medicine, 2017, 193, 130-139.	3.8	28
85	The genomic structure and developmental expression patterns of the human OPA-containing gene (HOPA). Human Genetics, 1999, 105, 174-178.	3.8	27
86	The DNA Methylation Signature of Smoking: An Archetype for the Identification of Biomarkers for Behavioral Illness. Nebraska Symposium on Motivation, 2014, 61, 109-127.	0.9	26
87	Ethnicity and Smoking-Associated DNA Methylation Changes at HIV Co-Receptor GPR15. Frontiers in Psychiatry, 2015, 6, 132.	2.6	26
88	Dihydropyridines modulate K+-evoked amino acid and adenosine release from cerebellar neuronal cultures. Neuroscience Letters, 1989, 102, 97-102.	2.1	25
89	Nonsupportive parenting affects telomere length in young adulthood among African Americans: Mediation through substance use Journal of Family Psychology, 2014, 28, 967-972.	1.3	25
90	Higher levels of protective parenting are associated with better young adult health: exploration of mediation through epigenetic influences on pro-inflammatory processes. Frontiers in Psychology, 2015, 6, 676.	2.1	25

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91	Developmental interplay between children's biobehavioral risk and the parenting environment from toddler to early school age: Prediction of socialization outcomes in preadolescence. Development and Psychopathology, 2015, 27, 775-790.	2.3	24
92	The association of the D2S2944 124 bp allele with recurrent early onset major depressive disorder in women. American Journal of Medical Genetics Part A, 2003, 121B, 39-43.	2.4	23
93	Transcriptional profiling of subjects from the Iowa adoption studies. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2007, 144B, 683-690.	1.7	23
94	Genomeâ€wide and digital polymerase chain reaction epigenetic assessments of alcohol consumption. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2018, 177, 479-488.	1.7	23
95	The Effect of Gender and Age at Onset of Depression on Mortality. Journal of Clinical Psychiatry, 1997, 58, 355-360.	2.2	23
96	Association of a D2S2944 allele with depression specifically among those with substance abuse or antisocial personality. Drug and Alcohol Dependence, 2006, 83, 33-41.	3.2	22
97	No association of the C677T methylenetetrahydrofolate reductase polymorphism with schizophrenia. Psychiatric Genetics, 2006, 16, 221-223.	1.1	22
98	DRD4 genotype moderates the impact of parental problems on unresolved loss or trauma. Attachment and Human Development, 2011, 13, 253-269.	2.1	22
99	Array-Based Epigenetic Aging Indices May Be Racially Biased. Genes, 2020, 11, 685.	2.4	22
100	K+- and temperature-evoked taurine efflux from hypothalamic astrocytes. Neuroscience Letters, 1990, 119, 23-26.	2.1	21
101	The relationship of the serotonin transporter (SLC6A4) extra long variant to gene expression in an African American sample. , 2012, 159B, 611-612.		21
102	Factors associated with sexual arousal, sexual sensation seeking and sexual satisfaction among female African American adolescents. Sexual Health, 2013, 10, 512.	0.9	21
103	Association of the HOPA12bp allele with a large X-chromosome haplotype and positive symptom schizophrenia. American Journal of Medical Genetics Part A, 2004, 127B, 20-27.	2.4	20
104	5-HTTLPR status moderates the effect of early adolescent substance use on risky sexual behavior Health Psychology, 2010, 29, 471-476.	1.6	20
105	MTHFR methylation moderates the impact of smoking on DNA methylation at AHRR for African American young adults. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2017, 174, 608-618.	1.7	20
106	The association of the HOPA12bp polymorphism with schizophrenia in the NIMH genetics initiative for schizophrenia sample. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2007, 144B, 743-747.	1.7	19
107	Examination of the Nicotine Dependence (NICSNP) Consortium findings in the lowa adoption studies population. Nicotine and Tobacco Research, 2009, 11, 286-292.	2.6	19
108	When inflammation and depression go together: The longitudinal effects of parent–child relationships. Development and Psychopathology, 2017, 29, 1969-1986.	2.3	19

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109	The Effect of Tobacco Smoking Differs across Indices of DNA Methylation-Based Aging in an African American Sample: DNA Methylation-Based Indices of Smoking Capture These Effects. Genes, 2020, 11, 311.	2.4	19
110	Transcriptional profiling of lymphoblast lines from subjects with panic disorder. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2007, 144B, 674-682.	1.7	18
111	Medical and psychiatric problems among men and women in a community corrections residential setting. Behavioral Sciences and the Law, 2009, 27, 695-711.	0.8	18
112	Stimulus-coupled taurine efflux from cerebellar neuronal cultures: On the roles of Ca++ and Na+. Journal of Neuroscience Research, 1989, 22, 167-171.	2.9	17
113	Genetic Moderation of the Impact of Parenting on Hostility Toward Romantic Partners. Journal of Marriage and Family, 2013, 75, 325-341.	2.6	17
114	The relationship of smoking to cg05575921 methylation in blood and saliva DNA samples from several studies. Scientific Reports, 2021, 11, 21627.	3.3	17
115	A meta-analysis of the association of the HOPA12bp polymorphism and schizophrenia. Psychiatric Genetics, 2006, 16, 73-76.	1.1	16
116	Financial strain, inflammatory factors, and haemoglobin <scp>A</scp> 1c levels in <scp>A</scp> frican <scp>A</scp> merican women. British Journal of Health Psychology, 2015, 20, 662-679.	3.5	16
117	A Direct Comparison of the Relationship of Epigenetic Aging and Epigenetic Substance Consumption Markers to Mortality in the Framingham Heart Study. Genes, 2019, 10, 51.	2.4	16
118	Saliva DNA Methylation Detects Nascent Smoking in Adolescents. Journal of Child and Adolescent Psychopharmacology, 2019, 29, 535-544.	1.3	16
119	A Four Marker Digital PCR Toolkit for Detecting Heavy Alcohol Consumption and the Effectiveness of Its Treatment. Journal of Insurance Medicine (New York, N Y), 2019, 48, 90-102.	0.2	16
120	Direct Sequencing of Trinucleotide Repeats from Cosmid Genomic DNA Template. Analytical Biochemistry, 1995, 225, 372-374.	2.4	15
121	The association of a HOPA polymorphism with major depression and phobia. Comprehensive Psychiatry, 2002, 43, 404-410.	3.1	15
122	A Cross-Platform Genome-Wide Comparison of the Relationship of Promoter DNA Methylation to Gene Expression. Frontiers in Genetics, 2012, 3, 12.	2.3	15
123	The relationship between alcohol consumption, perceived stress, and CRHR1 genotype on the hypothalamic–pituitary–adrenal axis in rural African Americans. Frontiers in Psychology, 2015, 6, 832.	2.1	15
124	Astrocyte Taurine. Annals of the New York Academy of Sciences, 1991, 633, 489-500.	3.8	14
125	Association of an exonicLDHA polymorphism with altered respiratory response in probands at high risk for panic disorder. American Journal of Medical Genetics Part A, 2003, 117B, 11-17.	2.4	14
126	Polymorphism analysis of HOPA: A candidate gene for schizophrenia. American Journal of Medical Genetics Part A, 2003, 123B, 33-38.	2.4	14

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127	Merging Genetic and Environmental Effects in the Iowa Adoption Studies: Focus on Depression. Annals of Clinical Psychiatry, 2006, 18, 219-222.	0.6	14
128	A pilot investigation of the impact of smoking cessation on biological age. American Journal on Addictions, 2017, 26, 129-135.	1.4	14
129	Testing Life Course Models Whereby Juvenile and Adult Adversity Combine to Influence Speed of Biological Aging. Journal of Health and Social Behavior, 2019, 60, 291-308.	4.8	14
130	Change in caregiver depression in response to parent training: Genetic moderation of intervention effects Journal of Family Psychology, 2009, 23, 112-117.	1.3	13
131	The Reversion of cg05575921 Methylation in Smoking Cessation: A Potential Tool for Incentivizing Healthy Aging. Genes, 2020, 11, 1415.	2.4	13
132	DNA methylation differentiates smoking from vaping and non-combustible tobacco use. Epigenetics, 2022, 17, 178-190.	2.7	13
133	Cigarette and Cannabis Smoking Effects on GPR15+ Helper T Cell Levels in Peripheral Blood: Relationships with Epigenetic Biomarkers. Genes, 2020, 11, 149.	2.4	13
134	Glutamate receptor agonists cause efflux of endogenous neuroactive amino acids from cerebellar neurons in culture. European Journal of Pharmacology, 1990, 177, 195-199.	3.5	12
135	K+-stimulated amino acid release from cultured cerebellar neurons: Comparison of static and dynamic stimulation paradigms. Neurochemical Research, 1991, 16, 899-904.	3.3	12
136	Looking Forward in Geriatric Anxiety and Depression: Implications of Basic Science for the Future. American Journal of Geriatric Psychiatry, 2005, 13, 1027-1040.	1.2	12
137	Interaction Between 5-HTTLPR Polymorphism and Abuse History on Adolescent African-American Females' Condom Use Behavior Following Participation in an HIV Prevention Intervention. Prevention Science, 2014, 15, 257-267.	2.6	12
138	Prevention of Early Substance Use Mediates, and Variation at SLC6A4 Moderates, SAAF Intervention Effects on OXTR Methylation. Prevention Science, 2018, 19, 90-100.	2.6	12
139	Refinement of cg05575921 demethylation response in nascent smoking. Clinical Epigenetics, 2020, 12, 92.	4.1	12
140	Perceived relationship support moderates the association of contextual stress with inflammation among African Americans Journal of Family Psychology, 2019, 33, 338-348.	1.3	12
141	The characterization and sequence analysis of thirty CTG-repeat containing genomic cosmid clones. European Journal of Human Genetics, 1998, 6, 89-94.	2.8	11
142	Exploring genetic moderators and epigenetic mediators of contextual and family effects: From Gene \tilde{A} — Environment to epigenetics. Development and Psychopathology, 2016, 28, 1333-1346.	2.3	11
143	Methylation of FKBP5 is associated with accelerated DNA methylation ageing and cardiometabolic risk: replication in young-adult and middle-aged Black Americans. Epigenetics, 2022, 17, 982-1002.	2.7	11
144	The inheritance of bipolar affective disorder: abundant genes coming together. Journal of Affective Disorders, 1997, 43, 1-3.	4.1	10

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145	A Review of Epigenetic Markers of Tobacco and Alcohol Consumption. Behavioral Sciences and the Law, 2015, 33, 675-690.	0.8	10
146	External validation of integrated genetic-epigenetic biomarkers for predicting incident coronary heart disease. Epigenomics, 2021, 13, 1095-1112.	2.1	10
147	AHRR Methylation is a Significant Predictor of Mortality Risk in Framingham Heart Study. Journal of Insurance Medicine (New York, N Y), 2019, 48, 79-89.	0.2	10
148	Childhood adversity predicts black young adults' DNA methylation-based accelerated aging: A dual pathway model. Development and Psychopathology, 2021, , 1-15.	2.3	10
149	Phorbol ester and dibutyryl cyclic AMP reduce content and efflux of taurine in primary cerebellar astrocytes in culture. Neuroscience Letters, 1988, 95, 323-328.	2.1	9
150	A Comparison of the Predictive Power of DNA Methylation with Carbohydrate Deficient Transferrin for Heavy Alcohol Consumption. Epigenetics, 2021, 16, 969-979.	2.7	9
151	Drs. Philibert and Carney-Doebbeling Reply. Psychosomatics, 2002, 43, 506.	2.5	8
152	An association study of PCQAP polymorphisms and schizophrenia. Psychiatric Genetics, 2004, 14, 169-172.	1.1	8
153	Childhood adversity is linked to adult health among African Americans via adolescent weight gain and effects are genetically moderated. Development and Psychopathology, 2021, 33, 803-820.	2.3	8
154	Role of elastin polymorphisms in panic disorder. American Journal of Medical Genetics Part A, 2003, 117B, 7-10.	2.4	7
155	Currents in Contemporary Ethics: Shocking Treatment: The Use of Tasers in Psychiatric Care. Journal of Law, Medicine and Ethics, 2006, 34, 116-120.	0.9	7
156	A Role for Epigenetics in Broadening the Scope of Pediatric Care in the Prevention of Adolescent Smoking. Epigenetic Diagnosis & Therapy, 2016, 1, 91-97.	0.1	7
157	Smoking in young adulthood among African Americans: Interconnected effects of supportive parenting in early adolescence, proinflammatory epitype, and young adult stress. Development and Psychopathology, 2017, 29, 957-969.	2.3	7
158	Unstable Childhood, Adult Adversity, and Smoking Accelerate Biological Aging Among Middle-Age African Americans: Similar Findings for GrimAge and PoAm. Journal of Aging and Health, 2022, 34, 487-498.	1.7	6
159	Reply to: Epstein-Barr Virus Transformed DNA as a Source of False Positive Findings in Methylation Studies of Psychiatric Conditions. Biological Psychiatry, 2011, 70, e27-e28.	1.3	5
160	Stress, relationship satisfaction, and health among African American women: Genetic moderation of effects Journal of Family Psychology, 2016, 30, 221-232.	1.3	5
161	Exon Array Biomarkers for the Differential Diagnosis of Schizophrenia and Bipolar Disorder. Molecular Neuropsychiatry, 2017, 3, 197-213.	2.9	5
162	Looking Forward in Geriatric Anxiety and Depression: Implications of Basic Science for the Future. American Journal of Geriatric Psychiatry, 2005, 13, 1027-1040.	1.2	5

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163	A Droplet Digital PCR Assay for Smoking Predicts All-Cause Mortality. Journal of Insurance Medicine (New York, N Y), 2018, 47, 220-229.	0.2	5
164	Shifts in lifestyle and socioeconomic circumstances predict change—for better or worse—in speed of epigenetic aging: A study of middle-aged black women. Social Science and Medicine, 2022, 307, 115175.	3.8	5
165	Inflammatory biomarker relationships with helper T cell GPR15 expression and cannabis and tobacco smoking. Journal of Psychosomatic Research, 2021, 141, 110326.	2.6	4
166	Epigenetic Analyses of Alcohol Consumption in Combustible and Non-Combustible Nicotine Product Users. Epigenomes, 2021, 5, 18.	1.8	4
167	Additive and Interactive Genetically Contextual Effects of HbA1c on cg19693031 Methylation in Type 2 Diabetes. Genes, 2022, 13, 683.	2.4	4
168	Therapeutic potential of targeting gene variants in schizophrenia. Expert Review of Neurotherapeutics, 2007, 7, 757-760.	2.8	3
169	Associations Between a Dopamine D4 Receptor Gene, Alcohol Use, and Sexual Behaviors Among Female Adolescent African Americans. Journal of HIV/AIDS and Social Services, 2015, 14, 136-153.	0.7	3
170	Methylation of MTHFR Moderates the Effect of Smoking on Genomewide Methylation Among Middle Age African Americans. Frontiers in Genetics, 2018, 9, 622.	2.3	3
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