

# Robert A Philibert

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1351819/publications.pdf>

Version: 2024-02-01

188  
papers

8,133  
citations

47006

47  
h-index

62596

80  
g-index

188  
all docs

188  
docs citations

188  
times ranked

8285  
citing authors

| #  | ARTICLE  | IF  | CITATIONS |
|----|--|-----|-----------|
| 1  | DNA methylation differentiates smoking from vaping and non-combustible tobacco use. <i>Epigenetics</i> , 2022, 17, 178-190.  | 2.7 | 13        |
| 2  | Methylation of FKBP5 is associated with accelerated DNA methylation ageing and cardiometabolic risk: replication in young-adult and middle-aged Black Americans. <i>Epigenetics</i> , 2022, 17, 982-1002.                        | 2.7 | 11        |
| 3  | Unstable Childhood, Adult Adversity, and Smoking Accelerate Biological Aging Among Middle-Age African Americans: Similar Findings for GrimAge and PoAm. <i>Journal of Aging and Health</i> , 2022, 34, 487-498.                  | 1.7 | 6         |
| 4  | Alcohol Use Intensity Decreases in Response to Successful Smoking Cessation Therapy. <i>Genes</i> , 2022, 13, 2.   | 2.4 | 2         |
| 5  | Additive and Interactive Genetically Contextual Effects of HbA1c on cg19693031 Methylation in Type 2 Diabetes. <i>Genes</i> , 2022, 13, 683.   | 2.4 | 4         |
| 6  | Shifts in lifestyle and socioeconomic circumstances predict changeâ€”for better or worseâ€”in speed of epigenetic aging: A study of middle-aged black women. <i>Social Science and Medicine</i> , 2022, 307, 115175.             | 3.8 | 5         |
| 7  | A Comparison of the Predictive Power of DNA Methylation with Carbohydrate Deficient Transferrin for Heavy Alcohol Consumption. <i>Epigenetics</i> , 2021, 16, 969-979.   | 2.7 | 9         |
| 8  | Childhood adversity is linked to adult health among African Americans via adolescent weight gain and effects are genetically moderated. <i>Development and Psychopathology</i> , 2021, 33, 803-820.                              | 2.3 | 8         |
| 9  | The effects of social adversity, discrimination, and health risk behaviors on the accelerated aging of African Americans: Further support for the weathering hypothesis. <i>Social Science and Medicine</i> , 2021, 282, 113169. | 3.8 | 98        |
| 10 | Inflammatory biomarker relationships with helper T cell GPR15 expression and cannabis and tobacco smoking. <i>Journal of Psychosomatic Research</i> , 2021, 141, 110326.   | 2.6 | 4         |
| 11 | The Reversion of DNA Methylation at Coronary Heart Disease Risk Loci in Response to Prevention Therapy. <i>Processes</i> , 2021, 9, 699.   | 2.8 | 3         |
| 12 | Costâ€”utility analysis of an integrated genetic/epigenetic test for assessing risk for coronary heart disease. <i>Epigenomics</i> , 2021, 13, 531-547.  | 2.1 | 2         |
| 13 | External validation of integrated genetic-epigenetic biomarkers for predicting incident coronary heart disease. <i>Epigenomics</i> , 2021, 13, 1095-1112.  | 2.1 | 10        |
| 14 | An Examination of Risk Factors for Tobacco and Cannabis Smoke Exposure in Adolescents Using an Epigenetic Biomarker. <i>Frontiers in Psychiatry</i> , 2021, 12, 688384.  | 2.6 | 3         |
| 15 | Epigenetic Analyses of Alcohol Consumption in Combustible and Non-Combustible Nicotine Product Users. <i>Epigenomes</i> , 2021, 5, 18.   | 1.8 | 4         |
| 16 | The relationship of smoking to cg05575921 methylation in blood and saliva DNA samples from several studies. <i>Scientific Reports</i> , 2021, 11, 21627.   | 3.3 | 17        |
| 17 | Childhood adversity predicts black young adultsâ€™ DNA methylation-based accelerated aging: A dual pathway model. <i>Development and Psychopathology</i> , 2021, , 1-15.   | 2.3 | 10        |
| 18 | AHRR methylation predicts smoking status and smoking intensity in both saliva and blood DNA. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2020, 183, 51-60.                                   | 1.7 | 55        |

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|----|---|-----|-----------|
| 19 | The Reversion of cg05575921 Methylation in Smoking Cessation: A Potential Tool for Incentivizing Healthy Aging. <i>Genes</i> , 2020, 11, 1415.  | 2.4 | 13        |
| 20 | A simple, rapid, interpretable, actionable and implementable digital PCR based mortality index. <i>Epigenetics</i> , 2020, 16, 1-15.  | 2.7 | 2         |
| 21 | 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. <i>Genes</i> , 2020, 11, 311.  | 2.4 | 19        |
| 22 | Array-Based Epigenetic Aging Indices May Be Racially Biased. <i>Genes</i> , 2020, 11, 685.  | 2.4 | 22        |
| 23 | Refinement of cg05575921 demethylation response in nascent smoking. <i>Clinical Epigenetics</i> , 2020, 12, 92.   | 4.1 | 12        |
| 24 | Cigarette and Cannabis Smoking Effects on GPR15+ Helper T Cell Levels in Peripheral Blood: Relationships with Epigenetic Biomarkers. <i>Genes</i> , 2020, 11, 149.  | 2.4 | 13        |
| 25 | Testing Life Course Models Whereby Juvenile and Adult Adversity Combine to Influence Speed of Biological Aging. <i>Journal of Health and Social Behavior</i> , 2019, 60, 291-308.   | 4.8 | 14        |
| 26 | A Direct Comparison of the Relationship of Epigenetic Aging and Epigenetic Substance Consumption Markers to Mortality in the Framingham Heart Study. <i>Genes</i> , 2019, 10, 51.   | 2.4 | 16        |
| 27 | Saliva DNA Methylation Detects Nascent Smoking in Adolescents. <i>Journal of Child and Adolescent Psychopharmacology</i> , 2019, 29, 535-544.   | 1.3 | 16        |
| 28 | Inflammation mediates the effect of discrimination, religiosity, and friendship network on expression of the Tp53 cancer suppressor gene. <i>SSM - Population Health</i> , 2019, 7, 100389.   | 2.7 | 2         |
| 29 | Neighborhood Disadvantage and Biological Aging: Using Marginal Structural Models to Assess the Link Between Neighborhood Census Variables and Epigenetic Aging. <i>Journals of Gerontology - Series B Psychological Sciences and Social Sciences</i> , 2019, 74, e50-e59. | 3.9 | 39        |
| 30 | Perceived relationship support moderates the association of contextual stress with inflammation among African Americans. <i>Journal of Family Psychology</i> , 2019, 33, 338-348.   | 1.3 | 12        |
| 31 | A Four Marker Digital PCR Toolkit for Detecting Heavy Alcohol Consumption and the Effectiveness of Its Treatment. <i>Journal of Insurance Medicine (New York, N Y)</i> , 2019, 48, 90-102.  | 0.2 | 16        |
| 32 | AHRR Methylation is a Significant Predictor of Mortality Risk in Framingham Heart Study. <i>Journal of Insurance Medicine (New York, N Y)</i> , 2019, 48, 79-89.  | 0.2 | 10        |
| 33 | Sharing the Burden of the Transition to Adulthood: African American Young Adults'™ Transition Challenges and Their Mothers'™ Health Risk. <i>American Sociological Review</i> , 2018, 83, 143-172.  | 5.2 | 40        |
| 34 | Genome-wide and digital polymerase chain reaction epigenetic assessments of alcohol consumption. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2018, 177, 479-488.  | 1.7 | 23        |
| 35 | Prevention of Early Substance Use Mediates, and Variation at SLC6A4 Moderates, SAAF Intervention Effects on OXTR Methylation. <i>Prevention Science</i> , 2018, 19, 90-100.   | 2.6 | 12        |
| 36 | Discrimination, segregation, and chronic inflammation: Testing the weathering explanation for the poor health of Black Americans. <i>Developmental Psychology</i> , 2018, 54, 1993-2006.  | 1.6 | 112       |

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|----|--|-----|-----------|
| 37 | Blood-Based Biomarkers for Predicting the Risk for Five-Year Incident Coronary Heart Disease in the Framingham Heart Study via Machine Learning. <i>Genes</i> , 2018, 9, 641.  | 2.4 | 29        |
| 38 | Methylation of MTHFR Moderates the Effect of Smoking on Genomewide Methylation Among Middle Age African Americans. <i>Frontiers in Genetics</i> , 2018, 9, 622.  | 2.3 | 3         |
| 39 | DNA methylation age is accelerated in alcohol dependence. <i>Translational Psychiatry</i> , 2018, 8, 182.  | 4.8 | 73        |
| 40 | MTHFR regulatory effects on methylation of CG05575921 in response to smoking: Effects are also discernable using MTHFR expression. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2018, 177, 529-534. | 1.7 | 1         |
| 41 | Would Addressing Alcohol Consumption Further Account for Variance in Methylation?. <i>American Journal of Psychiatry</i> , 2018, 175, 684-685.   | 7.2 | 2         |
| 42 | Dose Response and Prediction Characteristics of a Methylation Sensitive Digital PCR Assay for Cigarette Consumption in Adults. <i>Frontiers in Genetics</i> , 2018, 9, 137.  | 2.3 | 42        |
| 43 | Integrated genetic and epigenetic prediction of coronary heart disease in the Framingham Heart Study. <i>PLoS ONE</i> , 2018, 13, e0190549.  | 2.5 | 83        |
| 44 | A Droplet Digital PCR Assay for Smoking Predicts All-Cause Mortality. <i>Journal of Insurance Medicine (New York, N Y)</i> , 2018, 47, 220-229.  | 0.2 | 5         |
| 45 | A pilot investigation of the impact of smoking cessation on biological age. <i>American Journal on Addictions</i> , 2017, 26, 129-135.   | 1.4 | 14        |
| 46 | Smoking in young adulthood among African Americans: Interconnected effects of supportive parenting in early adolescence, proinflammatory epitype, and young adult stress. <i>Development and Psychopathology</i> , 2017, 29, 957-969.  | 2.3 | 7         |
| 47 | MTHFR methylation moderates the impact of smoking on DNA methylation at AHRR for African American young adults. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2017, 174, 608-618.                    | 1.7 | 20        |
| 48 | Accuracy and utility of an epigenetic biomarker for smoking in populations with varying rates of false self-report. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2017, 174, 641-650.                | 1.7 | 41        |
| 49 | An index of the ratio of inflammatory to antiviral cell types mediates the effects of social adversity and age on chronic illness. <i>Social Science and Medicine</i> , 2017, 185, 158-165.  | 3.8 | 31        |
| 50 | Methylation of the oxytocin receptor gene mediates the effect of adversity on negative schemas and depression. <i>Development and Psychopathology</i> , 2017, 29, 725-736.   | 2.3 | 44        |
| 51 | Childhood/Adolescent stressors and allostatic load in adulthood: Support for a calibration model. <i>Social Science and Medicine</i> , 2017, 193, 130-139.   | 3.8 | 28        |
| 52 | Genetically contextual effects of smoking on genome wide DNA methylation. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2017, 174, 595-607.  | 1.7 | 34        |
| 53 | Optimizing the chances of success in the search for epigenetic biomarkers: Embracing genetic variation. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2017, 174, 589-594.                            | 1.7 | 2         |
| 54 | When inflammation and depression go together: The longitudinal effects of parent-child relationships. <i>Development and Psychopathology</i> , 2017, 29, 1969-1986.  | 2.3 | 19        |

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|----|---|-----|-----------|
| 55 | Exon Array Biomarkers for the Differential Diagnosis of Schizophrenia and Bipolar Disorder. <i>Molecular Neuropsychiatry</i> , 2017, 3, 197-213.  | 2.9 | 5         |
| 56 | A Role for Epigenetics in Broadening the Scope of Pediatric Care in the Prevention of Adolescent Smoking. <i>Epigenetic Diagnosis &amp; Therapy</i> , 2016, 1, 91-97.   | 0.1 | 7         |
| 57 | Reversion of AHRR Demethylation Is a Quantitative Biomarker of Smoking Cessation. <i>Frontiers in Psychiatry</i> , 2016, 7, 55.   | 2.6 | 58        |
| 58 | Economic hardship and biological weathering: The epigenetics of aging in a U.S. sample of black women. <i>Social Science and Medicine</i> , 2016, 150, 192-200.   | 3.8 | 156       |
| 59 | A Contextual-Genetics Approach to Adolescent Drug Use and Sexual Risk Behavior. , 2016, , 399-426.  |     | 0         |
| 60 | Stress, relationship satisfaction, and health among African American women: Genetic moderation of effects.. <i>Journal of Family Psychology</i> , 2016, 30, 221-232.  | 1.3 | 5         |
| 61 | Exploring genetic moderators and epigenetic mediators of contextual and family effects: From Gene $\times$ Environment to epigenetics. <i>Development and Psychopathology</i> , 2016, 28, 1333-1346.  | 2.3 | 11        |
| 62 | Alcohol and tobacco consumption alter hypothalamic pituitary adrenal axis DNA methylation. <i>Psychoneuroendocrinology</i> , 2016, 66, 176-184.   | 2.7 | 33        |
| 63 | Parenting, Socioeconomic Status Risk, and Later Young Adult Health: Exploration of Opposing Indirect Effects via DNA Methylation. <i>Child Development</i> , 2016, 87, 111-121.   | 3.0 | 50        |
| 64 | 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. <i>Behavior Genetics</i> , 2016, 46, 151-169.                       | 2.1 | 98        |
| 65 | Developmental interplay between children's biobehavioral risk and the parenting environment from toddler to early school age: Prediction of socialization outcomes in preadolescence. <i>Development and Psychopathology</i> , 2015, 27, 775-790. | 2.3 | 24        |
| 66 | Methylomic Aging as a Window onto the Influence of Lifestyle: Tobacco and Alcohol Use Alter the Rate of Biological Aging. <i>Journal of the American Geriatrics Society</i> , 2015, 63, 2519-2525.  | 2.6 | 76        |
| 67 | Smoking, Methylation at AHRR, and Recidivism Risk in a Community Correction Sample of Individuals at High Risk for Recidivism. <i>Behavioral Sciences and the Law</i> , 2015, 33, 691-700.  | 0.8 | 1         |
| 68 | A Review of Epigenetic Markers of Tobacco and Alcohol Consumption. <i>Behavioral Sciences and the Law</i> , 2015, 33, 675-690.  | 0.8 | 10        |
| 69 | Financial strain, inflammatory factors, and haemoglobin $\times$ 1c levels in $\times$ frican $\times$ merican women. <i>British Journal of Health Psychology</i> , 2015, 20, 662-679.  | 3.5 | 16        |
| 70 | Current and Future Prospects for Epigenetic Biomarkers of Substance Use Disorders. <i>Genes</i> , 2015, 6, 991-1022.  | 2.4 | 70        |
| 71 | A quantitative epigenetic approach for the assessment of cigarette consumption. <i>Frontiers in Psychology</i> , 2015, 6, 656.  | 2.1 | 53        |
| 72 | Higher levels of protective parenting are associated with better young adult health: exploration of mediation through epigenetic influences on pro-inflammatory processes. <i>Frontiers in Psychology</i> , 2015, 6, 676.                         | 2.1 | 25        |

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|----|---|-----|-----------|
| 73 | Ethnicity and Smoking-Associated DNA Methylation Changes at HIV Co-Receptor GPR15. <i>Frontiers in Psychiatry</i> , 2015, 6, 132.   | 2.6 | 26        |
| 74 | The relationship between alcohol consumption, perceived stress, and CRHR1 genotype on the hypothalamic-pituitary-adrenal axis in rural African Americans. <i>Frontiers in Psychology</i> , 2015, 6, 832.                                    | 2.1 | 15        |
| 75 | Associations Between a Dopamine D4 Receptor Gene, Alcohol Use, and Sexual Behaviors Among Female Adolescent African Americans. <i>Journal of HIV/AIDS and Social Services</i> , 2015, 14, 136-153.  | 0.7 | 3         |
| 76 | Neighborhood crime and depressive symptoms among African American women: Genetic moderation and epigenetic mediation of effects. <i>Social Science and Medicine</i> , 2015, 146, 120-128.   | 3.8 | 47        |
| 77 | Prevention Effects Ameliorate the Prospective Association Between Nonsupportive Parenting and Diminished Telomere Length. <i>Prevention Science</i> , 2015, 16, 171-180.  | 2.6 | 48        |
| 78 | The genetic and epigenetic essentials of modern humans. , 2015, , 23-37.  |     | 2         |
| 79 | Nonsupportive parenting affects telomere length in young adulthood among African Americans: Mediation through substance use.. <i>Journal of Family Psychology</i> , 2014, 28, 967-972.  | 1.3 | 25        |
| 80 | The search for peripheral biomarkers for major depression: Benefiting from successes in the biology of smoking. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2014, 165, 230-234.                         | 1.7 | 1         |
| 81 | The DNA Methylation Signature of Smoking: An Archetype for the Identification of Biomarkers for Behavioral Illness. <i>Nebraska Symposium on Motivation</i> , 2014, 61, 109-127.  | 0.9 | 26        |
| 82 | A pilot examination of the genome-wide DNA methylation signatures of subjects entering and exiting short-term alcohol dependence treatment programs. <i>Epigenetics</i> , 2014, 9, 1212-1219.   | 2.7 | 72        |
| 83 | Is serotonin transporter genotype associated with epigenetic susceptibility or vulnerability? Examination of the impact of socioeconomic status risk on African American youth. <i>Development and Psychopathology</i> , 2014, 26, 289-304. | 2.3 | 41        |
| 84 | Methylation array data can simultaneously identify individuals and convey protected health information: an unrecognized ethical concern. <i>Clinical Epigenetics</i> , 2014, 6, 28.   | 4.1 | 35        |
| 85 | Interaction Between 5-HTTLPR Polymorphism and Abuse History on Adolescent African-American Females' Condom Use Behavior Following Participation in an HIV Prevention Intervention. <i>Prevention Science</i> , 2014, 15, 257-267.           | 2.6 | 12        |
| 86 | The effect of smoking on DNA methylation of peripheral blood mononuclear cells from African American women. <i>BMC Genomics</i> , 2014, 15, 151.  | 2.8 | 193       |
| 87 | 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. <i>Biological Psychology</i> , 2014, 96, 28-34.               | 2.2 | 31        |
| 88 | Epigenetic Silencing of the Human <i>NOS2</i> Gene: Rethinking the Role of Nitric Oxide in Human Macrophage Inflammatory Responses. <i>Journal of Immunology</i> , 2014, 192, 2326-2338.  | 0.8 | 107       |
| 89 | Harsh parenting and adolescent health: A longitudinal analysis with genetic moderation.. <i>Health Psychology</i> , 2014, 33, 401-409.  | 1.6 | 54        |
| 90 | Differential sensitivity to prevention programming: A dopaminergic polymorphism-enhanced prevention effect on protective parenting and adolescent substance use.. <i>Health Psychology</i> , 2014, 33, 182-191.                             | 1.6 | 55        |

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|-----|--|-----|-----------|
| 91  | Changes in DNA methylation at the aryl hydrocarbon receptor repressor may be a new biomarker for smoking. <i>Clinical Epigenetics</i> , 2013, 5, 19.   | 4.1 | 167       |
| 92  | Impact of child sex abuse on adult psychopathology: A genetically and epigenetically informed investigation.. <i>Journal of Family Psychology</i> , 2013, 27, 3-11.  | 1.3 | 52        |
| 93  | Supportive family environments, genes that confer sensitivity, and allostatic load among rural african american emerging adults: A prospective analysis.. <i>Journal of Family Psychology</i> , 2013, 27, 22-29.   | 1.3 | 39        |
| 94  | Genetic Moderation of the Impact of Parenting on Hostility Toward Romantic Partners. <i>Journal of Marriage and Family</i> , 2013, 75, 325-341.  | 2.6 | 17        |
| 95  | Cumulative socioeconomic status risk, allostatic load, and adjustment: A prospective latent profile analysis with contextual and genetic protective factors.. <i>Developmental Psychology</i> , 2013, 49, 913-927. | 1.6 | 112       |
| 96  | Factors associated with sexual arousal, sexual sensation seeking and sexual satisfaction among female African American adolescents. <i>Sexual Health</i> , 2013, 10, 512.  | 0.9 | 21        |
| 97  | Social Adversity, Genetic Variation, Street Code, and Aggression. <i>Youth Violence and Juvenile Justice</i> , 2012, 10, 3-24.   | 3.0 | 84        |
| 98  | Genetic moderation of contextual effects on negative arousal and parenting in African-American parents.. <i>Journal of Family Psychology</i> , 2012, 26, 46-55.  | 1.3 | 30        |
| 99  | Coordinated DNA methylation and gene expression changes in smoker alveolar macrophages: specific effects on VEGF receptor 1 expression. <i>Journal of Leukocyte Biology</i> , 2012, 92, 621-631.                   | 3.3 | 43        |
| 100 | Replication and meta-analysis of TMEM132D gene variants in panic disorder. <i>Translational Psychiatry</i> , 2012, 2, e156-e156.   | 4.8 | 74        |
| 101 | Life stress, the dopamine receptor gene, and emerging adult drug use trajectories: A longitudinal, multilevel, mediated moderation analysis. <i>Development and Psychopathology</i> , 2012, 24, 941-951.           | 2.3 | 38        |
| 102 | DNA CpG Methylation Contributes to Missing Nitric Oxide Inflammatory Response in Human Alveolar Macrophages. <i>Chest</i> , 2012, 142, 190A.   | 0.8 | 0         |
| 103 | The impact of stress on the life history strategies of African American adolescents: Cognitions, genetic moderation, and the role of discrimination.. <i>Developmental Psychology</i> , 2012, 48, 722-739.         | 1.6 | 74        |
| 104 | Demethylation of the aryl hydrocarbon receptor repressor as a biomarker for nascent smokers. <i>Epigenetics</i> , 2012, 7, 1331-1338.  | 2.7 | 146       |
| 105 | Effects of Genotype and Child Abuse on DNA Methylation and Gene Expression at the Serotonin Transporter. <i>Frontiers in Psychiatry</i> , 2012, 3, 55.   | 2.6 | 106       |
| 106 | A Cross-Platform Genome-Wide Comparison of the Relationship of Promoter DNA Methylation to Gene Expression. <i>Frontiers in Genetics</i> , 2012, 3, 12.  | 2.3 | 15        |
| 107 | The Impact of Recent Alcohol Use on Genome Wide DNA Methylation Signatures. <i>Frontiers in Genetics</i> , 2012, 3, 54.  | 2.3 | 110       |
| 108 | The relationship of the serotonin transporter (SLC6A4) extra long variant to gene expression in an African American sample. , 2012, 159B, 611-612.   |     | 21        |

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|-----|--|-----|-----------|
| 109 | Coordinated changes in AHRR methylation in lymphoblasts and pulmonary macrophages from smokers. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2012, 159B, 141-151.   | 1.7 | 230       |
| 110 | Reply to: Epstein-Barr Virus Transformed DNA as a Source of False Positive Findings in Methylation Studies of Psychiatric Conditions. <i>Biological Psychiatry</i> , 2011, 70, e27-e28.  | 1.3 | 5         |
| 111 | Gene environment interactions with a novel variable Monoamine Oxidase A transcriptional enhancer are associated with antisocial personality disorder. <i>Biological Psychology</i> , 2011, 87, 366-371.                                      | 2.2 | 40        |
| 112 | Methylation at 5HTT Mediates the Impact of Child Sex Abuse on Women's Antisocial Behavior: An Examination of the Iowa Adoptee Sample. <i>Psychosomatic Medicine</i> , 2011, 73, 83-87.   | 2.0 | 168       |
| 113 | Perceived discrimination, serotonin transporter linked polymorphic region status, and the development of conduct problems. <i>Development and Psychopathology</i> , 2011, 23, 617-627.   | 2.3 | 53        |
| 114 | Children's genotypes interact with maternal responsive care in predicting children's competence: Diathesisâ€“stress or differential susceptibility?. <i>Development and Psychopathology</i> , 2011, 23, 605-616.                             | 2.3 | 128       |
| 115 | The relationship of deiodinase 1 genotype and thyroid function to lifetime history of major depression in three independent populations. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2011, 156, 593-599. | 1.7 | 36        |
| 116 | Predictors of suicidal ideation, suicide attempts, and self-harm without lethal intent in a community corrections sample. <i>Journal of Criminal Justice</i> , 2011, 39, 238-245.  | 2.3 | 43        |
| 117 | DRD4 genotype moderates the impact of parental problems on unresolved loss or trauma. <i>Attachment and Human Development</i> , 2011, 13, 253-269.   | 2.1 | 22        |
| 118 | Social Environment, Genes, and Aggression. <i>American Sociological Review</i> , 2011, 76, 883-912.  | 5.2 | 160       |
| 119 | Child maltreatment moderates the association of MAOA with symptoms of depression and antisocial personality disorder.. <i>Journal of Family Psychology</i> , 2010, 24, 12-20.  | 1.3 | 95        |
| 120 | 5-HTTLPR status moderates the effect of early adolescent substance use on risky sexual behavior.. <i>Health Psychology</i> , 2010, 29, 471-476.  | 1.6 | 20        |
| 121 | Differential susceptibility to parenting among African American youths: Testing the DRD4 hypothesis.. <i>Journal of Family Psychology</i> , 2010, 24, 513-521.   | 1.3 | 78        |
| 122 | Methylation at <i>SLC6A4</i> is linked to family history of child abuse: An examination of the Iowa Adoptee sample. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2010, 153B, 710-713.                     | 1.7 | 209       |
| 123 | The effect of smoking on <i>MAOA</i> promoter methylation in DNA prepared from lymphoblasts and whole blood. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2010, 153B, 619-628.                            | 1.7 | 76        |
| 124 | Behavioral genetics in antisocial spectrum disorders and psychopathy: A review of the recent literature. <i>Behavioral Sciences and the Law</i> , 2010, 28, 148-173.   | 0.8 | 67        |
| 125 | Methylation Matters: Interaction Between Methylation Density and Serotonin Transporter Genotype Predicts Unresolved Loss or Trauma. <i>Biological Psychiatry</i> , 2010, 68, 405-407.  | 1.3 | 242       |
| 126 | Examination of the Nicotine Dependence (NICSNP) Consortium findings in the Iowa adoption studies population. <i>Nicotine and Tobacco Research</i> , 2009, 11, 286-292.   | 2.6 | 19        |



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|-----|--|-----|-----------|
| 127 | Participation in a Family-Centered Prevention Program Decreases Genetic Risk for Adolescents'™ Risky Behaviors. <i>Pediatrics</i> , 2009, 124, 911-917.  | 2.1 | 33        |
| 128 | Medical and psychiatric problems among men and women in a community corrections residential setting. <i>Behavioral Sciences and the Law</i> , 2009, 27, 695-711.   | 0.8 | 18        |
| 129 | Prevention Effects Moderate the Association of 5-HTTLPR and Youth Risk Behavior Initiation: Gene-Environment Hypotheses Tested via a Randomized Prevention Design. <i>Child Development</i> , 2009, 80, 645-661.   | 3.0 | 167       |
| 130 | Interplay of genes and early mother-child relationship in the development of self-regulation from toddler to preschool age. <i>Journal of Child Psychology and Psychiatry and Allied Disciplines</i> , 2009, 50, 1331-1338.  | 5.2 | 217       |
| 131 | Association between the serotonin transporter promoter polymorphism (5-HTTLPR) and adult unresolved attachment.. <i>Developmental Psychology</i> , 2009, 45, 64-76.  | 1.6 | 44        |
| 132 | Change in caregiver depression in response to parent training: Genetic moderation of intervention effects.. <i>Journal of Family Psychology</i> , 2009, 23, 112-117.   | 1.3 | 13        |
| 133 | Role of GABRA2 on risk for alcohol, nicotine, and cannabis dependence in the Iowa Adoption Studies. <i>Psychiatric Genetics</i> , 2009, 19, 91-98.   | 1.1 | 39        |
| 134 | Parenting moderates a genetic vulnerability factor in longitudinal increases in youths' substance use.. <i>Journal of Consulting and Clinical Psychology</i> , 2009, 77, 1-11.   | 2.0 | 102       |
| 135 | The relationship of 5HTT (<i>SLC6A4</i>) methylation and genotype on mRNA expression and liability to major depression and alcohol dependence in subjects from the Iowa Adoption Studies. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2008, 147B, 543-549. | 1.7 | 254       |
| 136 | MAOA methylation is associated with nicotine and alcohol dependence in women. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2008, 147B, 565-570.   | 1.7 | 142       |
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