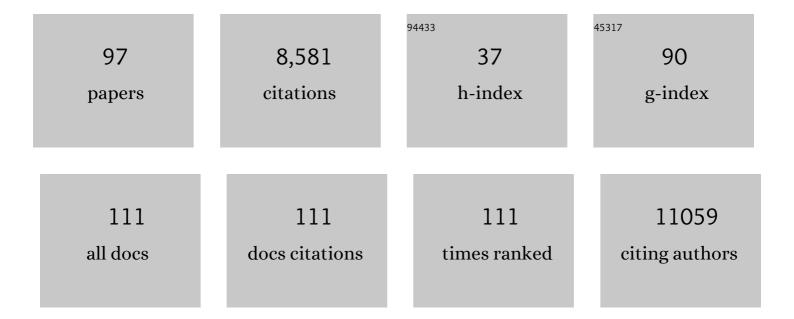


## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5493370/publications.pdf Version: 2024-02-01



KE XII

#	Article	IF	CITATIONS
1	A novel graph-based k-partitioning approach improves the detection of gene-gene correlations by single-cell RNA sequencing. BMC Genomics, 2022, 23, 35.	2.8	5
2	Epigenome-wide association study of posttraumatic stress disorder identifies novel loci in U.S. military veterans. Translational Psychiatry, 2022, 12, 65.	4.8	10
3	A new monocyte epigenetic clock reveals nonlinear effects of alcohol consumption on biological aging in three independent cohorts ( <i>N</i> Â=Â2242). Alcoholism: Clinical and Experimental Research, 2022, 46, 736-748.	2.4	9
4	Biological ageing with HIV infection: evaluating the geroscience hypothesis. The Lancet Healthy Longevity, 2022, 3, e194-e205.	4.6	20
5	Incorporating local ancestry improves identification of ancestry-associated methylation signatures and meQTLs in African Americans. Communications Biology, 2022, 5, 401.	4.4	3
6	The Stress and Resilience Town Hall: A systems response to support the health workforce during COVID-19 and beyond. General Hospital Psychiatry, 2022, 77, 80-87.	2.4	5
7	DNA methylation biomarker selected by an ensemble machine learning approach predicts mortality risk in an HIV-positive veteran population. Epigenetics, 2021, 16, 741-753.	2.7	9
8	DNA methylation signature on phosphatidylethanol, not onÂself-reported alcohol consumption, predicts hazardous alcohol consumption in two distinct populations. Molecular Psychiatry, 2021, 26, 2238-2253.	7.9	20
9	Design of the national adaptive trial for PTSD-related insomnia (NAP study), VA cooperative study program (CSP) #2016. Contemporary Clinical Trials, 2021, 109, 106540.	1.8	0
10	i2d: an R package for simulating data from images and the implications in biomedical research. Bioinformatics, 2021, 37, 2497-2498.	4.1	0
11	Psychological and biological resilience modulates the effects of stress on epigenetic aging. Translational Psychiatry, 2021, 11, 601.	4.8	44
12	Brush swab as a noninvasive surrogate for tissue biopsies in epigenomic profiling of oral cancer. Biomarker Research, 2021, 9, 90.	6.8	7
13	Epigenetic Associations With Estimated Clomerular Filtration Rate Among Men With Human Immunodeficiency Virus Infection. Clinical Infectious Diseases, 2020, 70, 667-673.	5.8	21
14	Chronic ketamine abuse is associated with orexin-A reduction and ACTH elevation. Psychopharmacology, 2020, 237, 45-53.	3.1	12
15	Association of Craving and Depressive Symptoms in Ketamineâ€Dependent Patients Undergoing Withdrawal Treatment. American Journal on Addictions, 2020, 29, 43-50.	1.4	20
16	Genome-wide association study of smoking trajectory and meta-analysis of smoking status in 842,000 individuals. Nature Communications, 2020, 11, 5302.	12.8	59
17	Comparison of methylation capture sequencing and Infinium MethylationEPIC array in peripheral blood mononuclear cells. Epigenetics and Chromatin, 2020, 13, 51.	3.9	26
18	DNA methylation mediates the effect of cocaine use on HIV severity. Clinical Epigenetics, 2020, 12, 140.	4.1	14

KE XU

#	Article	IF	CITATIONS
19	Searching for Genomic Biomarkers for Major Depressive Disorder in Peripheral Immune Cells. Biological Psychiatry, 2020, 88, 591-593.	1.3	1
20	Association of <i>OPRM1</i> Functional Coding Variant With Opioid Use Disorder. JAMA Psychiatry, 2020, 77, 1072.	11.0	135
21	Single-cell Transcriptome Mapping Identifies Common and Cell-type Specific Genes Affected by Acute Delta9-tetrahydrocannabinol in Humans. Scientific Reports, 2020, 10, 3450.	3.3	17
22	Gene selection for optimal prediction of cell position in tissues from single-cell transcriptomics data. Life Science Alliance, 2020, 3, e202000867.	2.8	20
23	Epigenomeâ€Wide <scp>DNA</scp> Methylation Association Analysis Identified Novel Loci in Peripheral Cells for Alcohol Consumption Among European American Male Veterans. Alcoholism: Clinical and Experimental Research, 2019, 43, 2111-2121.	2.4	27
24	Genome-wide association study of alcohol consumption and use disorder in 274,424 individuals from multiple populations. Nature Communications, 2019, 10, 1499.	12.8	346
25	Retrospective Association Analysis of Longitudinal Binary Traits Identifies Important Loci and Pathways in Cocaine Use. Genetics, 2019, 213, 1225-1236.	2.9	13
26	DNA Methylation Markers of Type 2 Diabetes Mellitus Among Male Veterans With or Without Human Immunodeficiency Virus Infection. Journal of Infectious Diseases, 2019, 219, 1959-1962.	4.0	20
27	Using DNA methylation to validate an electronic medical record phenotype for smoking. Addiction Biology, 2019, 24, 1056-1065.	2.6	11
28	Similar psychotic and cognitive profile between ketamine dependence with persistent psychosis and schizophrenia. Schizophrenia Research, 2018, 199, 313-318.	2.0	37
29	Cognitive profile of ketamine-dependent patients compared with methamphetamine-dependent patients and healthy controls. Psychopharmacology, 2018, 235, 2113-2121.	3.1	26
30	Epigenome-wide association analysis revealed that SOCS3 methylation influences the effect of cumulative stress on obesity. Biological Psychology, 2018, 131, 63-71.	2.2	49
31	Machine learning selected smoking-associated DNA methylation signatures that predict HIV prognosis and mortality. Clinical Epigenetics, 2018, 10, 155.	4.1	37
32	Decreased Blood Levels of Oxytocin in Ketamine-Dependent Patients During Early Abstinence. Frontiers in Psychiatry, 2018, 9, 633.	2.6	14
33	The profile of cognitive impairments in chronic ketamine users. Psychiatry Research, 2018, 266, 124-131.	3.3	44
34	AUDIT  and ICD codes as phenotypes for harmful alcohol use: association with <i>ADH1B</i> polymorphisms in two US populations. Addiction, 2018, 113, 2214-2224.	3.3	24
35	Validating Harmful Alcohol Use as a Phenotype for Genetic Discovery Using Phosphatidylethanol and a Polymorphism in <i><scp>ADH</scp>1B</i> . Alcoholism: Clinical and Experimental Research, 2017, 41, 998-1003.	2.4	15
36	Alcohol Triggers Reemergence of Ketamine-Like Experience in a Ketamine Ex-User. Journal of Clinical Psychopharmacology, 2017, 37, 110-112.	1.4	3

#	Article	IF	CITATIONS
37	Identification of HIV infection-related DNA methylation sites and advanced epigenetic aging in HIV-positive, treatment-naive U.S. veterans. Aids, 2017, 31, 571-575.	2.2	49
38	Longitudinal SNPâ€set association analysis of quantitative phenotypes. Genetic Epidemiology, 2017, 41, 81-93.	1.3	13
39	DNA methylation signatures of illicit drug injection and hepatitis C are associated with HIV frailty. Nature Communications, 2017, 8, 2243.	12.8	32
40	Epigenome-wide differential DNA methylation between HIV-infected and uninfected individuals. Epigenetics, 2016, 11, 750-760.	2.7	78
41	Profiling the psychotic, depressive and anxiety symptoms in chronic ketamine users. Psychiatry Research, 2016, 237, 311-315.	3.3	37
42	Genomewide Association Study for Maximum Number of Alcoholic Drinks in European Americans and African Americans. Alcoholism: Clinical and Experimental Research, 2015, 39, 1137-1147.	2.4	58
43	The Effects of BDNF Val66Met Gene Polymorphism on Serum BDNF and Cognitive Function in Methamphetamine-Dependent Patients and Normal Controls. Journal of Clinical Psychopharmacology, 2015, 35, 517-524.	1.4	20
44	Effect of Tricuspid Regurgitation and the Right Heart on Survival After Transcatheter Aortic Valve Replacement. Circulation: Cardiovascular Interventions, 2015, 8, .	3.9	148
45	Relationship of serum levels of TNF-α, IL-6 and IL-18 and schizophrenia-like symptoms in chronic ketamine abusers. Schizophrenia Research, 2015, 169, 10-15.	2.0	13
46	Preliminary analysis of positive and negative syndrome scale in ketamine-associated psychosis in comparison with schizophrenia. Journal of Psychiatric Research, 2015, 61, 64-72.	3.1	50
47	Serum level of vascular endothelial growth factor decreased in chronic ketamine abusers. Drug and Alcohol Dependence, 2015, 152, 57-61.	3.2	8
48	The analysis of BDNF gene polymorphism haplotypes and impulsivity in methamphetamine abusers. Comprehensive Psychiatry, 2015, 59, 62-67.	3.1	11
49	Repeated ketamine administration alters <i>N</i> -methyl- <scp>d</scp> -aspartic acid receptor subunit gene expression: Implication of genetic vulnerability for ketamine abuse and ketamine psychosis in humans. Experimental Biology and Medicine, 2015, 240, 145-155.	2.4	46
50	Development of a checklist of short-term and long-term psychological symptoms associated with ketamine use. Shanghai Archives of Psychiatry, 2015, 27, 186-94.	0.7	5
51	Targeting KRAS mutant non-small cell lung cancer (NSCLC) with deltarasin: A small molecule inhibitor of KRAS-PDEI ´ interaction Journal of Clinical Oncology, 2015, 33, e13597-e13597.	1.6	1
52	Impact of Aortic Annulus Size on Valve Hemodynamics and Clinical Outcomes After Transcatheter and Surgical Aortic Valve Replacement. Circulation: Cardiovascular Interventions, 2014, 7, 701-711.	3.9	90
53	The Joint Effects of <i><scp>ADH</scp>1B</i> Variants and Childhood Adversity on Alcohol Related Phenotypes in Africanâ€American and Europeanâ€American Women and Men. Alcoholism: Clinical and Experimental Research, 2014, 38, 2907-2914.	2.4	28
54	Serum brain-derived neurotrophic factor and nerve growth factor decreased in chronic ketamine abusers. Drug and Alcohol Dependence, 2014, 142, 290-294.	3.2	30

#	Article	IF	CITATIONS
55	An association between BDNF Val66Met polymorphism and impulsivity in methamphetamine abusers. Neuroscience Letters, 2014, 582, 16-20.	2.1	22
56	Improvement of quality of life in methadone treatment patients in northern Taiwan: a follow-up study. BMC Psychiatry, 2013, 13, 190.	2.6	34
57	A variant on the kappa opioid receptor gene (OPRK1) is associated with stress response and related drug craving, limbic brain activation and cocaine relapse risk. Translational Psychiatry, 2013, 3, e292-e292.	4.8	49
58	Rater Evaluations for Psychiatric Instruments and Cultural Differences. Journal of Nervous and Mental Disease, 2012, 200, 814-820.	1.0	6
59	Cognitive and serum BDNF correlates of BDNF Val66Met gene polymorphism in patients with schizophrenia and normal controls. Human Genetics, 2012, 131, 1187-1195.	3.8	103
60	Novelty seeking involved in mediating the association between COMT gene and onset age of drug use in Chinese heroin-dependent patients. Neuroscience Research, 2011, 71, e398.	1.9	0
61	Comparing the PANSS in Chinese and American inpatients: Cross-cultural psychiatric analyses of instrument translation and implementation. Schizophrenia Research, 2011, 132, 146-152.	2.0	22
62	Role of Novelty Seeking Personality Traits as Mediator of the Association between COMT and Onset Age of Drug Use in Chinese Heroin Dependent Patients. PLoS ONE, 2011, 6, e22923.	2.5	19
63	Cognitive effects of genetic variation in monoamine neurotransmitter systems: A populationâ€based study of <i>COMT</i> , <i>MAOA</i> , and <i>5HTTLPR</i> . American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2011, 156, 158-167.	1.7	41
64	Effects of Catechol- <i>O</i> -Methyltransferase on Normal Variation in the Cognitive Function of Children. American Journal of Psychiatry, 2009, 166, 909-916.	7.2	61
65	Emotional symptoms in children: The effect of maternal depression, life events, and COMT genotype. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2009, 150B, 209-218.	1.7	21
66	Zhou et al. reply. Nature, 2009, 458, E7-E7.	27.8	1
67	Genetic variation in human NPY expression affects stress response and emotion. Nature, 2008, 452, 997-1001.	27.8	387
68	BHLHB2 Controls <i>Bdnf</i> Promoter 4 Activity and Neuronal Excitability. Journal of Neuroscience, 2008, 28, 1118-1130.	3.6	83
69	Association between ADORA2A and DRD2 Polymorphisms and Caffeine-Induced Anxiety. Neuropsychopharmacology, 2008, 33, 2791-2800.	5.4	209
70	Addictions Biology: Haplotype-Based Analysis for 130 Candidate Genes on a Single Array. Alcohol and Alcoholism, 2008, 43, 505-515.	1.6	222
71	Gender-Specific Effects of the Catechol- <i>O</i> -Methyltransferase <i>Val</i> <sup>108/158</sup> <i>Met</i> Polymorphism on Cognitive Function in Children. American Journal of Psychiatry, 2007, 164, 142-149.	7.2	115
72	Irritable assault and variation in the COMT gene. Psychiatric Genetics, 2007, 17, 344-346.	1.1	13

#	Article	IF	CITATIONS
73	Diplotypes of the Human Serotonin 1B Receptor Promoter Predict Growth Hormone Responses to Sumatriptan in Abstinent Alcohol-Dependent Men. Biological Psychiatry, 2007, 61, 974-978.	1.3	7
74	Minor differences in haplotype frequency estimates can produce very large differences in heterogeneity test statistics. BMC Genetics, 2007, 8, 38.	2.7	12
75	Using ancestry-informative markers to define populations and detect population stratification. Journal of Psychopharmacology, 2006, 20, 19-26.	4.0	115
76	Serotonin Transporter Promoter Gain-of-Function Genotypes Are Linked to Obsessive-Compulsive Disorder. American Journal of Human Genetics, 2006, 78, 815-826.	6.2	1,032
77	Catechol-O-methyltransferase Val158Met genotype variation is associated with prefrontal-dependent task performance in schizotypal personality disorder patients and comparison groups. Psychiatric Genetics, 2006, 16, 117-124.	1.1	40
78	Imaging Genomics Applied to Anxiety, Stress Response, and Resiliency. Neuroinformatics, 2006, 4, 51-64.	2.8	31
79	DRD2 Promoter Region Variation as a Predictor of Sustained Response to Antipsychotic Medication in First-Episode Schizophrenia Patients. American Journal of Psychiatry, 2006, 163, 529-531.	7.2	146
80	Estimation of Haplotypes at DRD2 May Have Produced Misleading Results—Reply. Archives of General Psychiatry, 2006, 63, 939.	12.3	0
81	Association of COMT Val158Met Genotype With Executive Functioning Following Traumatic Brain Injury. Journal of Neuropsychiatry and Clinical Neurosciences, 2005, 17, 465-471.	1.8	90
82	Genomics and variation of ionotropic glutamate receptors: implications for neuroplasticity. Amino Acids, 2005, 28, 169-175.	2.7	7
83	Genetic basis for individual variations in pain perception and the development of a chronic pain condition. Human Molecular Genetics, 2005, 14, 135-143.	2.9	1,134
84	BDNF Variation and Mood Disorders: A Novel Functional Promoter Polymorphism and Val66Met are Associated with Anxiety but Have Opposing Effects. Neuropsychopharmacology, 2005, 30, 1353-1361.	5.4	223
85	Association of Specific Haplotypes of D2 Dopamine ReceptorGene With Vulnerability to Heroin Dependence in 2 Distinct Populations. Archives of General Psychiatry, 2004, 61, 597.	12.3	119
86	Differential expression of human COMT alleles in brain and lymphoblasts detected by RT-coupled 5? nuclease assay. Psychopharmacology, 2004, 177, 178-184.	3.1	55
87	Genetic influence on variability in human acute experimental pain sensitivity associated with gender, ethnicity and psychological temperament. Pain, 2004, 109, 488-496.	4.2	355
88	Association of the G1947A COMT (Val108/158Met) gene polymorphism with prefrontal P300 during information processing. Biological Psychiatry, 2003, 54, 40-48.	1.3	160
89	COMT <i> val <sup>158</sup> met </i> Genotype Affects µ-Opioid Neurotransmitter Responses to a Pain Stressor. Science, 2003, 299, 1240-1243.	12.6	1,025
90	Genetic origins of anxiety in women: a role for a functional catechol-O-methyltransferase polymorphism. Psychiatric Genetics, 2003, 13, 33-41.	1.1	250

#	Article	IF	CITATIONS
91	Relationship of the delta-opioid receptor gene to heroin abuse in a large Chinese case/control sample. American Journal of Medical Genetics Part A, 2002, 110, 45-50.	2.4	46
92	Immune related genetic polymorphisms and schizophrenia among the Chinese. Human Immunology, 2001, 62, 714-724.	2.4	32
93	DNA Melting Analysis for Detection of Single Nucleotide Polymorphisms. Clinical Chemistry, 2001, 47, 635-644.	3.2	92
94	Nuclear factor l°B is a critical determinant in N-methyl-d-aspartate receptor-mediated neuroprotection. Journal of Neurochemistry, 2001, 78, 254-264.	3.9	155
95	Catechol-O-methyltransferase Vall58Met polymorphism: frequency analysis in Han Chinese subjects and allelic association of the low activity allele with bipolar affective disorder. Pharmacogenetics and Genomics, 1997, 7, 349-353.	5.7	93
96	DNA methylation signatures analysis with Illumina Infinitum MethylationEPIC and Infinium Human Methylation 450K BeadChip. Protocol Exchange, 0, , .	0.3	1
97	Functional alleles, neuroimaging and intermediate phenotypes in the deconstruction of complex behavioral variation. , 0, , 365-382.		0