

Genome-wide gene-environment interaction in depression candidate genes

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Citation Report

#	ARTICLE	IF	CITATIONS
1	FKBP5 polymorphisms and hypothalamic-pituitary-adrenal axis negative feedback in major depression and obsessive-compulsive disorder. <i>Journal of Psychiatric Research</i> , 2018, 104, 227-234.	1.5	19
2	Machine learning in major depression: From classification to treatment outcome prediction. <i>CNS Neuroscience and Therapeutics</i> , 2018, 24, 1037-1052.	1.9	199
3	Non-coding genetic variation shaping mental health. <i>Current Opinion in Psychology</i> , 2019, 27, 18-24.	2.5	14
4	Association of childhood traumatization and neuropsychiatric outcomes with altered plasma micro RNA-levels. <i>Neuropsychopharmacology</i> , 2019, 44, 2030-2037.	2.8	21
5	Genomics and psychological resilience: a research agenda. <i>Molecular Psychiatry</i> , 2019, 24, 1770-1778.	4.1	54
6	The effect of a genetic variant at the schizophrenia associated AS3MT/BORCS7 locus on striatal dopamine function: A PET imaging study. <i>Psychiatry Research - Neuroimaging</i> , 2019, 291, 34-41.	0.9	13
7	Long-term environmental impact on object recognition, spatial memory, and reversal learning capabilities in <i>Cacna1c</i> haploinsufficient rats. <i>Human Molecular Genetics</i> , 2019, 28, 4113-4131.	1.4	9
8	A Fast and Accurate Method for Genome-wide Scale Phenome-wide G – E Analysis and Its Application to UK Biobank. <i>American Journal of Human Genetics</i> , 2019, 105, 1182-1192.	2.6	20
9	Epigenome-wide association study of depression symptomatology in elderly monozygotic twins. <i>Translational Psychiatry</i> , 2019, 9, 214.	2.4	48
10	No Support for Historical Candidate Gene or Candidate Gene-by-Interaction Hypotheses for Major Depression Across Multiple Large Samples. <i>American Journal of Psychiatry</i> , 2019, 176, 376-387.	4.0	436
11	What DNA methylation modifications and/or genetic variations interact with childhood maltreatment in the development of depression: A systematic review. <i>Journal of Affective Disorders</i> , 2019, 252, 325-333.	2.0	11
12	Gene-Environment Interactions and Epigenetic Mechanisms in Depression. , 2019, , 17-25.		2
13	Genetic variants in major depressive disorder: From pathophysiology to therapy. , 2019, 194, 22-43.		57
14	Continuity of genetic and environmental influences on clinically assessed major depression from ages 18 to 45. <i>Psychological Medicine</i> , 2019, 49, 2582-2590.	2.7	5
15	Suicide and suicide behaviors: A review of transcriptomics and multiomics studies in psychiatric disorders. <i>Journal of Neuroscience Research</i> , 2020, 98, 601-615.	1.3	14
16	Genetics of resilience: Implications from genome-wide association studies and candidate genes of the stress response system in posttraumatic stress disorder and depression. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2020, 183, 77-94.	1.1	54
17	Gene-environment interaction in psychiatry. , 2020, , 363-373.		0
18	AVPR1A main effect and OXTR-by-environment interplay in individual differences in depression level. <i>Heliyon</i> , 2020, 6, e05240.	1.4	6

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19	Interactions of childhood maltreatment and genetic variations in adult depression: A systematic review. <i>Journal of Affective Disorders</i> , 2020, 276, 119-136.	2.0	9
20	Genome-wide association studies of antidepressant class response and treatment-resistant depression. <i>Translational Psychiatry</i> , 2020, 10, 360.	2.4	33
21	A systematic review of gene-by-intervention studies of alcohol and other substance use. <i>Development and Psychopathology</i> , 2021, 33, 1410-1427.	1.4	6
22	A Summary of Recent Updates on the Genetic Determinants of Depression. , 2020, , 1-27.		1
23	Interaction between serotonin and dopamine and impulsivity: A gene-gene - interaction approach. <i>Personality and Individual Differences</i> , 2021, 169, 110014.	1.6	2
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25	Polygenic Risk for Major Depression Interacts with Parental Criticism in Predicting Adolescent Depressive Symptom Development. <i>Journal of Youth and Adolescence</i> , 2021, 50, 159-176.	1.9	14
26	Progress in Neurophysiological Mechanisms of Depression. <i>Advances in Psychology</i> , 2021, 11, 1759-1767.	0.0	0
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30	Cumulative procedural pain and brain development in very preterm infants: A systematic review of clinical and preclinical studies. <i>Neuroscience and Biobehavioral Reviews</i> , 2021, 123, 320-336.	2.9	20
31	Associations and interactions of the serotonin receptor genes 5-HT1A, 5-HT2A, and childhood trauma with alexithymia in two independent general-population samples. <i>Psychiatry Research</i> , 2021, 298, 113783.	1.7	6
32	Sex-stratified gene-by-environment genome-wide interaction study of trauma, posttraumatic-stress, and suicidality. <i>Neurobiology of Stress</i> , 2021, 14, 100309.	1.9	32
33	The associations of <i>CNR1</i> SNPs and haplotypes with vulnerability and treatment response phenotypes in Han Chinese with major depressive disorder: A case-control association study. <i>Molecular Genetics & Genomic Medicine</i> , 2021, 9, e1752.	0.6	6
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36	Genetic Architecture of Depression: Where Do We Stand Now?. <i>Advances in Experimental Medicine and Biology</i> , 2021, 1305, 203-230.	0.8	4

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37	Transcriptional and Epigenetic Regulation of the Corticotropin-Releasing Hormone System and Genetic Associations With Neuropsychiatric Disorders. , 2021, , 83-94.		1
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42	Genetic Advance in Depressive Disorder. Advances in Experimental Medicine and Biology, 2019, 1180, 19-57.	0.8	2
43	<p>Geneâ“Environment Interactions in Major Mental Disorders in the Czech Republic</p>. Neuropsychiatric Disease and Treatment, 2020, Volume 16, 1147-1156.	1.0	0
44	Unipolar depression. , 2020, , 613-631.		0
45	Proteomic profiling of postmortem prefrontal cortex tissue of suicide completers. Translational Psychiatry, 2022, 12, 142.	2.4	3
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48	Air pollution, genetic factors and the risk of depression. Science of the Total Environment, 2022, 850, 158001.	3.9	15
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50	New Research Perspectives on the Interplay Between Genes and Environment on Executive Function Development. Biological Psychiatry, 2023, 94, 131-141.	0.7	7
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52	Genetic risk of depression is different in subgroups of dietary ratio of tryptophan to large neutral amino acids. Scientific Reports, 2023, 13, .	1.6	0
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