Gerard van Grootheest

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/9264594/publications.pdf

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43 papers

6,054 citations

236612 25 h-index 42 g-index

46 all docs 46 docs citations

46 times ranked

12640 citing authors

#	Article	IF	CITATIONS
1	Genetic relationship between five psychiatric disorders estimated from genome-wide SNPs. Nature Genetics, 2013, 45, 984-994.	9.4	2,067
2	A mega-analysis of genome-wide association studies for major depressive disorder. Molecular Psychiatry, 2013, 18, 497-511.	4.1	1,002
3	Psychiatric genome-wide association study analyses implicate neuronal, immune and histone pathways. Nature Neuroscience, 2015, 18, 199-209.	7.1	701
4	Heritability and genomics of gene expression in peripheral blood. Nature Genetics, 2014, 46, 430-437.	9.4	370
5	Joint Analysis of Psychiatric Disorders Increases Accuracy of Risk Prediction for Schizophrenia, Bipolar Disorder, and Major Depressive Disorder. American Journal of Human Genetics, 2015, 96, 283-294.	2.6	225
6	Gene expression in major depressive disorder. Molecular Psychiatry, 2016, 21, 339-347.	4.1	178
7	Genome-wide physical activity interactions in adiposity ― A meta-analysis of 200,452 adults. PLoS Genetics, 2017, 13, e1006528.	1.5	158
8	Sex differences in the human peripheral blood transcriptome. BMC Genomics, 2014, 15, 33.	1.2	131
9	Harmonization of Neuroticism and Extraversion phenotypes across inventories and cohorts in the Genetics of Personality Consortium: an application of Item Response Theory. Behavior Genetics, 2014, 44, 295-313.	1.4	103
10	Effect of Multinutrient Supplementation and Food-Related Behavioral Activation Therapy on Prevention of Major Depressive Disorder Among Overweight or Obese Adults With Subsyndromal Depressive Symptoms. JAMA - Journal of the American Medical Association, 2019, 321, 858.	3.8	88
11	An Analysis of Two Genome-wide Association Meta-analyses Identifies a New Locus for Broad Depression Phenotype. Biological Psychiatry, 2017, 82, 322-329.	0.7	84
12	Differential involvement of the dorsal hippocampus in passive avoidance in C57bl/6J and DBA/2J mice. Hippocampus, 2008, 18 , $11-19$.	0.9	78
13	Genetic effects influencing risk for major depressive disorder in China and Europe. Translational Psychiatry, 2017, 7, e1074-e1074.	2.4	64
14	Applying polygenic risk scores to postpartum depression. Archives of Women's Mental Health, 2014, 17, 519-528.	1.2	62
15	Differential gene expression patterns between smokers and nonâ€smokers: cause or consequence?. Addiction Biology, 2017, 22, 550-560.	1.4	62
16	Methylome-wide association findings for major depressive disorder overlap in blood and brain and replicate in independent brain samples. Molecular Psychiatry, 2020, 25, 1344-1354.	4.1	61
17	The association between lower educational attainment and depression owing to shared genetic effects? Results in ~25 000 subjects. Molecular Psychiatry, 2015, 20, 735-743.	4.1	59
18	A methylation study of long-term depression risk. Molecular Psychiatry, 2020, 25, 1334-1343.	4.1	56

#	Article	IF	Citations
19	Genome-wide analyses of borderline personality features. Molecular Psychiatry, 2014, 19, 923-929.	4.1	55
20	Prevention of depression through nutritional strategies in high-risk persons: rationale and design of the MooDFOOD prevention trial. BMC Psychiatry, 2016, 16, 192.	1.1	52
21	Integrating evolutionary and regulatory information with a multispecies approach implicates genes and pathways in obsessive-compulsive disorder. Nature Communications, 2017, 8, 774.	5.8	52
22	HIV-infected mental health patients: characteristics and comparison with HIV-infected patients from the general population and non-infected mental health patients. BMC Psychiatry, 2013, 13, 35.	1.1	50
23	Huntingtin gene repeat size variations affect risk of lifetime depression. Translational Psychiatry, 2017, 7, 1277.	2.4	37
24	Cell Type–Specific Methylome-wide Association Studies Implicate Neurotrophin and Innate Immune Signaling in Major Depressive Disorder. Biological Psychiatry, 2020, 87, 431-442.	0.7	35
25	Biomarker-based subtyping of depression and anxiety disorders using Latent Class Analysis. A NESDA study. Psychological Medicine, 2019, 49, 617-627.	2.7	27
26	Correcting for cell-type effects in DNA methylation studies: reference-based method outperforms latent variable approaches in empirical studies. Genome Biology, 2017, 18, 24.	3.8	25
27	Large normal-range TBP and ATXN7 CAG repeat lengths are associated with increased lifetime risk of depression. Translational Psychiatry, 2017, 7, e1143-e1143.	2.4	20
28	Working memory moderates the relation between the brain-derived neurotropic factor (BDNF) and psychotherapy outcome for depression. Journal of Psychiatric Research, 2020, 130, 424-432.	1.5	17
29	Convergence of evidence from a methylome-wide CpG-SNP association study and GWAS of major depressive disorder. Translational Psychiatry, 2018, 8, 162.	2.4	16
30	Effect of food-related behavioral activation therapy on food intake and the environmental impact of the diet: results from the MooDFOOD prevention trial. European Journal of Nutrition, 2020, 59, 2579-2591.	1.8	15
31	Gene transcripts associated with muscle strength: a CHARGE meta-analysis of 7,781 persons. Physiological Genomics, 2016, 48, 1-11.	1.0	11
32	Depressive Symptom Clusters in Relation to Body Weight Status: Results From Two Large European Multicenter Studies. Frontiers in Psychiatry, 2019, 10, 858.	1.3	11
33	Novel neuronal surface autoantibodies in plasma of patients with depression and anxiety. Translational Psychiatry, 2020, 10, 404.	2.4	10
34	Nutrition and depression: Summary of findings from the EUâ€funded MooDFOOD depression prevention randomised controlled trial and a critical review of the literature. Nutrition Bulletin, 2020, 45, 403-414.	0.8	8
35	Fat metabolism is associated with telomere length in six population-based studies. Human Molecular Genetics, 2022, 31, 1159-1170.	1.4	7
36	Determinants of plasma 25-hydroxyvitamin D levels in healthy adults in the Netherlands. Netherlands Journal of Medicine, 2014, 72, 533-40.	0.6	7

#	Article	IF	CITATIONS
37	Acceptability and feasibility of two interventions in the MooDFOOD Trial: a food-related depression prevention randomised controlled trial in overweight adults with subsyndromal symptoms of depression. BMJ Open, 2020, 10, e034025.	0.8	4
38	Associations of carotid intima media thickness with gene expression in whole blood and genetically predicted gene expression across 48 tissues. Human Molecular Genetics, 2022, 31, 1171-1182.	1.4	4
39	Habitual Behavior as a Mediator Between Food-Related Behavioral Activation and Change in Symptoms of Depression in the MooDFOOD Trial. Clinical Psychological Science, 2021, 9, 649-665.	2.4	4
40	Toxoplasma gondii seropositivity in patients with depressive and anxiety disorders. Brain, Behavior, & Immunity - Health, 2021, 11, 100197.	1.3	3
41	PREDICTING THE FUTURE DISEASE STATUS OF DEPRESSED PATIENTS FROM DNA METHYLATION PATTERNS IN BLOOD. European Neuropsychopharmacology, 2019, 29, S793-S794.	0.3	O
42	Overweight and obese individuals with depressive symptoms from the MooDFOOD prevention trial: Role of sociodemographic, somatic health, and weight related factors. Journal of Affective Disorders Reports, 2021, 4, 100126.	0.9	0
43	Toxoplasma gondii seropositivity in patients with depressive and anxiety disorders. European Psychiatry, 2021, 64, S75-S75.	0.1	0