

# Maren Lang

## List of Publications by Year in descending order

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Version: 2024-02-01

35  
papers

3,266  
citations

430754

18  
h-index

377752

34  
g-index

36  
all docs

36  
docs citations

36  
times ranked

6245  
citing authors

#	ARTICLE	IF	CITATIONS
1	Lifetime and current depression in the German National Cohort (NAKO). <i>World Journal of Biological Psychiatry</i> , 2023, 24, 865-880.	1.3	18
2	Interaction Testing and Polygenic Risk Scoring to Estimate the Association of Common Genetic Variants With Treatment Resistance in Schizophrenia. <i>JAMA Psychiatry</i> , 2022, 79, 260.	6.0	44
3	Urbanicity, behavior problems and HPA axis regulation in preschoolers. <i>Psychoneuroendocrinology</i> , 2022, 137, 105660.	1.3	1
4	Meta-analysis of epigenome-wide associations between DNA methylation at birth and childhood cognitive skills. <i>Molecular Psychiatry</i> , 2022, 27, 2126-2135.	4.1	13
5	Mapping genomic loci implicates genes and synaptic biology in schizophrenia. <i>Nature</i> , 2022, 604, 502-508.	13.7	929
6	DNA methylation signatures of aggression and closely related constructs: A meta-analysis of epigenome-wide studies across the lifespan. <i>Molecular Psychiatry</i> , 2021, 26, 2148-2162.	4.1	21
7	“The Heidelberg Five” personality dimensions: Genome-wide associations, polygenic risk for neuroticism, and psychopathology 20 years after assessment. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2021, 186, 77-89.	1.1	6
8	Methylome-wide change associated with response to electroconvulsive therapy in depressed patients. <i>Translational Psychiatry</i> , 2021, 11, 347.	2.4	12
9	Microbiome profiles are associated with cognitive functioning in 45-month-old children. <i>Brain, Behavior, and Immunity</i> , 2021, 98, 151-160.	2.0	18
10	Association of Locomotor Activity During Sleep Deprivation Treatment With Response. <i>Frontiers in Psychiatry</i> , 2020, 11, 688.	1.3	2
11	Attitudes toward the right to autonomous decision-making in psychiatric genetic testing: Controversial and context-dependent. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2019, 180, 555-565.	1.1	6
12	Longitudinal transcriptome-wide gene expression analysis of sleep deprivation treatment shows involvement of circadian genes and immune pathways. <i>Translational Psychiatry</i> , 2019, 9, 343.	2.4	21
13	Evidence for increased genetic risk load for major depression in patients assigned to electroconvulsive therapy. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2019, 180, 35-45.	1.1	18
14	Efficient region-based test strategy uncovers genetic risk factors for functional outcome in bipolar disorder. <i>European Neuropsychopharmacology</i> , 2019, 29, 156-170.	0.3	7
15	Association of the alcohol dehydrogenase gene polymorphism rs1789891 with gray matter brain volume, alcohol consumption, alcohol craving and relapse risk. <i>Addiction Biology</i> , 2019, 24, 110-120.	1.4	13
16	Impact on birth weight of maternal smoking throughout pregnancy mediated by DNA methylation. <i>BMC Genomics</i> , 2018, 19, 290.	1.2	41
17	Response to therapeutic sleep deprivation: a naturalistic study of clinical and genetic factors and post-treatment depressive symptom trajectory. <i>Neuropsychopharmacology</i> , 2018, 43, 2572-2577.	2.8	17
18	Shared genetic etiology between alcohol dependence and major depressive disorder. <i>Psychiatric Genetics</i> , 2018, 28, 66-70.	0.6	19

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19	Telomere Length in Newborns is Related to Maternal Stress During Pregnancy. <i>Neuropsychopharmacology</i> , 2017, 42, 2407-2413.	2.8	83
20	Investigation of <i>SHANK3</i> in schizophrenia. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2017, 174, 390-398.	1.1	34
21	Associations between SNPs and immune-related circulating proteins in schizophrenia. <i>Scientific Reports</i> , 2017, 7, 12586.	1.6	21
22	Hair Cortisol in Twins: Heritability and Genetic Overlap with Psychological Variables and Stress-System Genes. <i>Scientific Reports</i> , 2017, 7, 15351.	1.6	50
23	Contribution of copy number variants to schizophrenia from a genome-wide study of 41,321 subjects. <i>Nature Genetics</i> , 2017, 49, 27-35.	9.4	838
24	Hair Cortisol and Its Association With Psychological Risk Factors for Psychiatric Disorders: A Pilot Study in Adolescent Twins. <i>Twin Research and Human Genetics</i> , 2016, 19, 438-446.	0.3	31
25	Perceived stress and hair cortisol: Differences in bipolar disorder and schizophrenia. <i>Psychoneuroendocrinology</i> , 2016, 69, 26-34.	1.3	48
26	Genome-wide association study of 40,000 individuals identifies two novel loci associated with bipolar disorder. <i>Human Molecular Genetics</i> , 2016, 25, 3383-3394.	1.4	182
27	No Reliable Association between Runs of Homozygosity and Schizophrenia in a Well-Powered Replication Study. <i>PLoS Genetics</i> , 2016, 12, e1006343.	1.5	24
28	The effects of single nucleotide polymorphisms in glutamatergic neurotransmission genes on neural response to alcohol cues and craving. <i>Addiction Biology</i> , 2015, 20, 1022-1032.	1.4	30
29	Increased mesolimbic cue-reactivity in carriers of the mu-opioid-receptor gene OPRM1 A118G polymorphism predicts drinking outcome: A functional imaging study in alcohol dependent subjects. <i>European Neuropsychopharmacology</i> , 2015, 25, 1128-1135.	0.3	46
30	New-born females show higher stress- and genotype-independent methylation of SLC6A4 than males. <i>Borderline Personality Disorder and Emotion Dysregulation</i> , 2015, 2, 8.	1.1	38
31	XRCC5 as a Risk Gene for Alcohol Dependence: Evidence from a Genome-Wide Gene-Set-Based Analysis and Follow-up Studies in Drosophila and Humans. <i>Neuropsychopharmacology</i> , 2015, 40, 361-371.	2.8	12
32	A genome-wide association study confirms PNPLA3 and identifies TM6SF2 and MBOAT7 as risk loci for alcohol-related cirrhosis. <i>Nature Genetics</i> , 2015, 47, 1443-1448.	9.4	435
33	Investigation of the involvement of <i>MIR185</i> and its target genes in the development of schizophrenia. <i>Journal of Psychiatry and Neuroscience</i> , 2014, 39, 386-396.	1.4	23
34	ADH1B Arg48His Allele Frequency Map: Filling in the Gap for Central Europe. <i>Biological Psychiatry</i> , 2014, 75, e15.	0.7	11
35	Genome-wide significant association between alcohol dependence and a variant in the <i>ADH</i> gene cluster. <i>Addiction Biology</i> , 2012, 17, 171-180.	1.4	154