

# Cynthia Marie-Claire

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3488020/publications.pdf>

Version: 2024-02-01

60  
papers

2,046  
citations

257450

24  
h-index

265206

42  
g-index

61  
all docs

61  
docs citations

61  
times ranked

3092  
citing authors

#	ARTICLE	IF	CITATIONS
1	Neurobiological and behavioral mechanisms of circadian rhythm disruption in bipolar disorder: A critical multi-disciplinary literature review and agenda for future research from the ISBD task force on chronobiology. <i>Bipolar Disorders</i> , 2022, 24, 232-263.	1.9	36
2	Occurrence and severity of cocaine-induced hallucinations: Two distinct phenotypes with shared clinical factors but specific genetic risk factors. <i>Drug and Alcohol Dependence</i> , 2022, 232, 109270.	3.2	2
3	Influence of childhood maltreatment on prevalence, onset, and persistence of psychiatric comorbidities and suicide attempts in bipolar disorders. <i>European Psychiatry</i> , 2022, 65, 1-32.	0.2	8
4	Methylomic Biomarkers of Lithium Response in Bipolar Disorder: A Proof of Transferability Study. <i>Pharmaceuticals</i> , 2022, 15, 133.	3.8	5
5	Association between childhood maltreatment and the clinical course of bipolar disorders: A survival analysis of mood recurrences. <i>Acta Psychiatrica Scandinavica</i> , 2022, 145, 373-383.	4.5	9
6	The molecular pathophysiology of mood disorders: From the analysis of single molecular layers to multi-omic integration. <i>Progress in Neuro-Psychopharmacology and Biological Psychiatry</i> , 2022, 116, 110520.	4.8	6
7	Telomere length and mitochondrial DNA copy number in bipolar disorder: identification of a subgroup of young individuals with accelerated cellular aging. <i>Translational Psychiatry</i> , 2022, 12, 135.	4.8	15
8	Clustering suicidal phenotypes and genetic associations with brain-derived neurotrophic factor in patients with substance use disorders. <i>Translational Psychiatry</i> , 2021, 11, 72.	4.8	4
9	Biomarkers to predict staging and treatment response in opioid dependence: A narrative review. <i>Drug Development Research</i> , 2021, 82, 668-677.	2.9	1
10	Clinical Trials of Cannabidiol for Substance Use Disorders: Outcome Measures, Surrogate Endpoints, and Biomarkers. <i>Frontiers in Psychiatry</i> , 2021, 12, 565617.	2.6	5
11	Network of co-expressed circadian genes, childhood maltreatment and sleep quality in bipolar disorders. <i>Chronobiology International</i> , 2021, 38, 986-993.	2.0	7
12	Mini review: Recent advances on epigenetic effects of lithium. <i>Neuroscience Letters</i> , 2021, 761, 136116.	2.1	5
13	A Comparison of Different Approaches to Clinical Phenotyping of Lithium Response: A Proof of Principle Study Employing Genetic Variants of Three Candidate Circadian Genes. <i>Pharmaceuticals</i> , 2021, 14, 1072.	3.8	2
14	Combining schizophrenia and depression polygenic risk scores improves the genetic prediction of lithium response in bipolar disorder patients. <i>Translational Psychiatry</i> , 2021, 11, 606.	4.8	25
15	Lithium effects on serine-threonine kinases activity: High throughput kinomic profiling of lymphoblastoid cell lines from excellent-responders and non-responders bipolar patients. <i>World Journal of Biological Psychiatry</i> , 2020, 21, 317-324.	2.6	4
16	Childhood maltreatment and HPA axis gene expression in bipolar disorders: A gene network analysis. <i>Psychoneuroendocrinology</i> , 2020, 120, 104753.	2.7	6
17	A DNA methylation signature discriminates between excellent and non-response to lithium in patients with bipolar disorder type 1. <i>Scientific Reports</i> , 2020, 10, 12239.	3.3	21
18	Translational study of the whole transcriptome in rats and genetic polymorphisms in humans identifies LRP1B and VPS13A as key genes involved in tolerance to cocaine-induced motor disturbances. <i>Translational Psychiatry</i> , 2020, 10, 381.	4.8	6

#	ARTICLE	IF	CITATIONS
19	QT length during methadone maintenance treatment: gene–dose interaction. <i>Fundamental and Clinical Pharmacology</i> , 2019, 33, 96-106.	1.9	10
20	Determination of sets of covariating gene expression using graph analysis on pairwise expression ratios. <i>Bioinformatics</i> , 2019, 35, 258-265.	4.1	9
21	Selecting reference genes in RT-qPCR based on equivalence tests: a network based approach. <i>Scientific Reports</i> , 2019, 9, 16231.	3.3	15
22	Increased plasma levels of high mobility group box 1 protein in patients with bipolar disorder: A pilot study. <i>Journal of Neuroimmunology</i> , 2019, 334, 576993.	2.3	5
23	Dopamine ( DRD 2 ) and Serotonin ( HTR 2A, 2C ) Receptor Gene Polymorphisms do not influence early response to Risperidone in South Indian Patients with Schizophrenia. <i>Fundamental and Clinical Pharmacology</i> , 2019, 33, 355-364.	1.9	11
24	A tutorial on conducting genome-wide association studies: Quality control and statistical analysis. <i>International Journal of Methods in Psychiatric Research</i> , 2018, 27, e1608.	2.1	465
25	Lithium response in bipolar disorders and core clock genes expression. <i>World Journal of Biological Psychiatry</i> , 2018, 19, 619-632.	2.6	45
26	DNA Methylation as a Biomarker of Treatment Response Variability in Serious Mental Illnesses: A Systematic Review Focused on Bipolar Disorder, Schizophrenia, and Major Depressive Disorder. <i>International Journal of Molecular Sciences</i> , 2018, 19, 3026.	4.1	38
27	Molecular Signatures of Lithium Treatment: Current Knowledge. <i>Pharmacopsychiatry</i> , 2018, 51, 212-219.	3.3	18
28	Analysis of the Influence of microRNAs in Lithium Response in Bipolar Disorder. <i>Frontiers in Psychiatry</i> , 2018, 9, 207.	2.6	28
29	Lithium response in bipolar disorder: No difference in GADL1 gene expression between cell lines from excellent-responders and non-responders. <i>Psychiatry Research</i> , 2017, 251, 217-220.	3.3	9
30	Pharmacoeugenomics of opiates and methadone maintenance treatment: current data and perspectives. <i>Pharmacogenomics</i> , 2017, 18, 1359-1372.	1.3	8
31	Lithium Response Variability: New Avenues and Hypotheses. , 2017, , 157-178.		2
32	Circadian genes and lithium response in bipolar disorders: associations with <i>PPARGC1A</i> ( <i>PGC-1<math>\alpha</math></i> ) and <i>RORA</i> . <i>Genes, Brain and Behavior</i> , 2016, 15, 660-668.	2.2	37
33	Variability of response to methadone: genome-wide DNA methylation analysis in two independent cohorts. <i>Epigenomics</i> , 2016, 8, 181-195.	2.1	17
34	Impact of P-glycoprotein at the blood-brain barrier on the uptake of heroin and its main metabolites: behavioral effects and consequences on the transcriptional responses and reinforcing properties. <i>Psychopharmacology</i> , 2014, 231, 3139-3149.	3.1	30
35	Comparison of the transcriptional responses induced by acute morphine, methadone and buprenorphine. <i>European Journal of Pharmacology</i> , 2013, 711, 10-18.	3.5	8
36	Transport of Biogenic Amine Neurotransmitters at the Mouse Blood–Retina and Blood–Brain Barriers by Uptake1 and Uptake2. <i>Journal of Cerebral Blood Flow and Metabolism</i> , 2012, 32, 1989-2001.	4.3	34

#	ARTICLE	IF	CITATIONS
37	Chronic and intermittent morphine treatment differently regulates opioid and dopamine systems: a role in locomotor sensitization. <i>Psychopharmacology</i> , 2011, 216, 297-303.	3.1	39
38	PRECLINICAL STUDY: Modulation of MDMA-induced behavioral and transcriptional effects by the delta opioid antagonist naltrindole in mice. <i>Addiction Biology</i> , 2009, 14, 245-252.	2.6	14
39	Effect of chronic exposure to morphine on the rat blood-brain barrier: focus on the P-glycoprotein. <i>Journal of Neurochemistry</i> , 2008, 107, 647-657.	3.9	60
40	Involvement of D1 dopamine receptor in MDMA-induced locomotor activity and striatal gene expression in mice. <i>Brain Research</i> , 2008, 1211, 1-5.	2.2	43
41	Characteristics of dual specificity phosphatases mRNA regulation by 3,4-methylenedioxymethamphetamine acute treatment in mice striatum. <i>Brain Research</i> , 2008, 1239, 42-48.	2.2	14
42	Effects of the selective neurotensin antagonist SR 142948A on 3,4-methylenedioxymethamphetamine-induced behaviours in mice. <i>Neuropharmacology</i> , 2008, 54, 1107-1111.	4.1	14
43	Effects of chronic morphine and morphine withdrawal on gene expression in rat peripheral blood mononuclear cells. <i>Neuropharmacology</i> , 2008, 55, 1347-1354.	4.1	32
44	Sensitization to the conditioned rewarding effects of morphine modulates gene expression in rat hippocampus. <i>Neuropharmacology</i> , 2007, 52, 430-435.	4.1	16
45	Rnd family genes are differentially regulated by 3,4-methylenedioxymethamphetamine and cocaine acute treatment in mice brain. <i>Brain Research</i> , 2007, 1134, 12-17.	2.2	29
46	Expression of drug transporters at the blood-brain barrier using an optimized isolated rat brain microvessel strategy. <i>Brain Research</i> , 2007, 1134, 1-11.	2.2	125
47	Regulation of genes involved in dopamine transporter modulation by acute cocaine in rat striatum. <i>Neuroscience Letters</i> , 2006, 398, 235-240.	2.1	11
48	Analysis of transcriptional responses in the mouse dorsal striatum following acute 3,4-methylenedioxymethamphetamine (ecstasy): Identification of extracellular signal-regulated kinase-controlled genes. <i>Neuroscience</i> , 2006, 137, 473-482.	2.3	17
49	Cytoskeletal Genes Regulation by Chronic Morphine Treatment in Rat Striatum. <i>Neuropsychopharmacology</i> , 2004, 29, 2208-2215.	5.4	86
50	Further evidence that the CCK <sub>2</sub> receptor is coupled to two transduction pathways using site-directed mutagenesis. <i>Journal of Neurochemistry</i> , 2003, 85, 454-461.	3.9	25
51	Importance of ERK activation in behavioral and biochemical effects induced by MDMA in mice. <i>British Journal of Pharmacology</i> , 2003, 140, 831-838.	5.4	111
52	General function of N-terminal propeptide on assisting protein folding and inhibiting catalytic activity based on observations with a chimeric thermolysin-like protease. <i>Biochemical and Biophysical Research Communications</i> , 2003, 301, 1093-1098.	2.1	38
53	Fos but not Cart (cocaine and amphetamine regulated transcript) is overexpressed by several drugs of abuse: a comparative study using real-time quantitative polymerase chain reaction in rat brain. <i>Neuroscience Letters</i> , 2003, 345, 77-80.	2.1	62
54	Folding pathway mediated by an intramolecular chaperone: the structural and functional characterization of the aqualysin I propeptide. <i>Journal of Molecular Biology</i> , 2001, 305, 151-165.	4.2	46

#	ARTICLE	IF	CITATIONS
55	Exploration of the S1 subsite of neprilysin: A joined molecular modeling and site-directed mutagenesis study. , 2000, 39, 365-371.		11
56	The prosequence of thermolysin acts as an intramolecular chaperone when expressed in trans with the mature sequence in Escherichia coli 1 Edited by A. R. Fersht. Journal of Molecular Biology, 1999, 285, 1911-1915.	4.2	62
57	Differences in transition state stabilization between thermolysin (EC 3.4.24.27) and neprilysin (EC Tj ETQq1 1 0.784314 rgBT /Overl	2.8	28
58	Characterization of Glu350 as a Critical Residue Involved in the N-Terminal Amine Binding Site of Aminopeptidase N (EC 3.4.11.2): Insights into Its Mechanism of Action. Biochemistry, 1998, 37, 686-692.	2.5	111
59	Intramolecular Processing of Prothermolysin. Journal of Biological Chemistry, 1998, 273, 5697-5701.	3.4	44
60	Evidence by Site-Directed Mutagenesis That Arginine 203 of Thermolysin and Arginine 717 of Neprilysin (Neutral Endopeptidase) Play Equivalent Critical Roles in Substrate Hydrolysis and Inhibitor Binding. Biochemistry, 1997, 36, 13938-13945.	2.5	41