

Marti Jett

List of Publications by Year in descending order

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31
papers

1,345
citations

489802

18
h-index

488211

31
g-index

34
all docs

34
docs citations

34
times ranked

2578
citing authors

#	ARTICLE	IF	CITATIONS
1	Enhancing Discovery of Genetic Variants for Posttraumatic Stress Disorder Through Integration of Quantitative Phenotypes and Trauma Exposure Information. <i>Biological Psychiatry</i> , 2022, 91, 626-636.	0.7	21
2	Acute and Delayed Effects of Stress Eliciting Post-Traumatic Stress-Like Disorder Differentially Alters Fecal Microbiota Composition in a Male Mouse Model. <i>Frontiers in Cellular and Infection Microbiology</i> , 2022, 12, 810815.	1.8	6
3	Transcriptomics of Wet Skin Biopsies Predict Early Radiation-Induced Hematological Damage in a Mouse Model. <i>Genes</i> , 2022, 13, 538.	1.0	0
4	Epigenetic biotypes of post-traumatic stress disorder in war-zone exposed veteran and active duty males. <i>Molecular Psychiatry</i> , 2021, 26, 4300-4314.	4.1	22
5	A DNA methylation clock associated with age-related illnesses and mortality is accelerated in men with combat PTSD. <i>Molecular Psychiatry</i> , 2021, 26, 4999-5009.	4.1	52
6	Pre-deployment risk factors for PTSD in active-duty personnel deployed to Afghanistan: a machine-learning approach for analyzing multivariate predictors. <i>Molecular Psychiatry</i> , 2021, 26, 5011-5022.	4.1	55
7	PTSD is associated with increased DNA methylation across regions of HLA-DPB1 and SPATC1L. <i>Brain, Behavior, and Immunity</i> , 2021, 91, 429-436.	2.0	17
8	TBI weight-drop model with variable impact heights differentially perturbs hippocampus-cerebellum specific transcriptomic profile. <i>Experimental Neurology</i> , 2021, 335, 113516.	2.0	11
9	Utilization of machine learning for identifying symptom severity military-related PTSD subtypes and their biological correlates. <i>Translational Psychiatry</i> , 2021, 11, 227.	2.4	11
10	â€œGrimAge,â€•an epigenetic predictor of mortality, is accelerated in major depressive disorder. <i>Translational Psychiatry</i> , 2021, 11, 193.	2.4	46
11	Longitudinal genome-wide methylation study of PTSD treatment using prolonged exposure and hydrocortisone. <i>Translational Psychiatry</i> , 2021, 11, 398.	2.4	14
12	Diallyl Sulfide Attenuation of Carcinogenesis in Mammary Epithelial Cells through the Inhibition of ROS Formation, and DNA Strand Breaks. <i>Biomolecules</i> , 2021, 11, 1313.	1.8	3
13	Multi-omic biomarker identification and validation for diagnosing warzone-related post-traumatic stress disorder. <i>Molecular Psychiatry</i> , 2020, 25, 3337-3349.	4.1	68
14	Molecular genetic overlap between posttraumatic stress disorder and sleep phenotypes. <i>Sleep</i> , 2020, 43, .	0.6	32
15	Evaluating the impact of trauma and PTSD on epigenetic prediction of lifespan and neural integrity. <i>Neuropsychopharmacology</i> , 2020, 45, 1609-1616.	2.8	63
16	Novel Pharmacological Targets for Combat PTSDâ€™Metabolism, Inflammation, The Gut Microbiome, and Mitochondrial Dysfunction. <i>Military Medicine</i> , 2020, 185, 311-318.	0.4	24
17	Genomic influences on self-reported childhood maltreatment. <i>Translational Psychiatry</i> , 2020, 10, 38.	2.4	47
18	Effect of Combat Exposure and Posttraumatic Stress Disorder on Telomere Length and Amygdala Volume. <i>Biological Psychiatry: Cognitive Neuroscience and Neuroimaging</i> , 2020, 5, 678-687.	1.1	10

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19	Predeployment neurocognitive functioning predicts postdeployment posttraumatic stress in Army personnel.. <i>Neuropsychology</i> , 2020, 34, 276-287.	1.0	22
20	Mechanistic inferences on metabolic dysfunction in posttraumatic stress disorder from an integrated model and multiomic analysis: role of glucocorticoid receptor sensitivity. <i>American Journal of Physiology - Endocrinology and Metabolism</i> , 2019, 317, E879-E898.	1.8	22
21	International meta-analysis of PTSD genome-wide association studies identifies sex- and ancestry-specific genetic risk loci. <i>Nature Communications</i> , 2019, 10, 4558.	5.8	363
22	Polygenic risk associated with post-traumatic stress disorder onset and severity. <i>Translational Psychiatry</i> , 2019, 9, 165.	2.4	23
23	Molecular alterations induced by <i>Yersinia pestis</i> , dengue virus and Staphylococcal enterotoxin B under severe stress. <i>Brain, Behavior, and Immunity</i> , 2019, 80, 725-741.	2.0	2
24	Metabolomic analysis of male combat veterans with post traumatic stress disorder. <i>PLoS ONE</i> , 2019, 14, e0213839.	1.1	54
25	Metabolism, Metabolomics, and Inflammation in Posttraumatic Stress Disorder. <i>Biological Psychiatry</i> , 2018, 83, 866-875.	0.7	131
26	Epigenetic Age in Male Combat-Exposed War Veterans: Associations with Posttraumatic Stress Disorder Status. <i>Molecular Neuropsychiatry</i> , 2018, 4, 90-99.	3.0	35
27	Systems biology approach to understanding post-traumatic stress disorder. <i>Molecular BioSystems</i> , 2015, 11, 980-993.	2.9	20
28	Acute and Chronic Plasma Metabolomic and Liver Transcriptomic Stress Effects in a Mouse Model with Features of Post-Traumatic Stress Disorder. <i>PLoS ONE</i> , 2015, 10, e0117092.	1.1	36
29	Molecular evidence of stress-induced acute heart injury in a mouse model simulating posttraumatic stress disorder. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 3188-3193.	3.3	45
30	Conserved MHC Class II-Presented Dengue Virus Epitopes Identified by Immunoproteomics Analysis Are Targets for Cross-Serotype Reactive T-Cell Response. <i>Journal of Infectious Diseases</i> , 2012, 205, 647-655.	1.9	31
31	Murine model of repeated exposures to conspecific trained aggressors simulates features of post-traumatic stress disorder. <i>Behavioural Brain Research</i> , 2012, 235, 55-66.	1.2	46