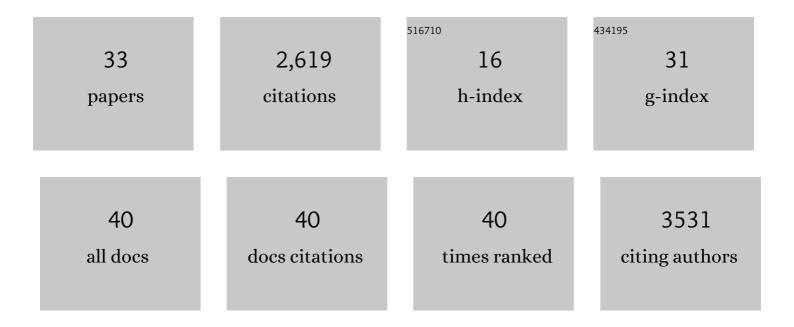
Lisa M Tarantino

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/2386901/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	The Collaborative Cross, a community resource for the genetic analysis of complex traits. Nature Genetics, 2004, 36, 1133-1137.	21.4	1,034
2	The Intestinal Microbiota in Acute Anorexia Nervosa and During Renourishment. Psychosomatic Medicine, 2015, 77, 969-981.	2.0	237
3	Analyses of allele-specific gene expression in highly divergent mouse crosses identifies pervasive allelic imbalance. Nature Genetics, 2015, 47, 353-360.	21.4	204
4	High genetic susceptibility to ethanol withdrawal predicts low ethanol consumption. Mammalian Genome, 1998, 9, 983-990.	2.2	152
5	Behavior and mutagenesis screens: the importance of baseline analysis of inbred strains. Mammalian Genome, 2000, 11, 555-564.	2.2	151
6	Confirmation of Quantitative Trait Loci for Alcohol Preference in Mice. Alcoholism: Clinical and Experimental Research, 1998, 22, 1099-1105.	2.4	123
7	A Common and Unstable Copy Number Variant Is Associated with Differences in Glo1 Expression and Anxiety-Like Behavior. PLoS ONE, 2009, 4, e4649.	2.5	108
8	Quantitative-trait loci analysis of cocaine-related behaviours and neurochemistry. Pharmacogenetics and Genomics, 1999, 9, 607-618.	5.7	81
9	Maternal vitamin D depletion alters DNA methylation at imprinted loci in multiple generations. Clinical Epigenetics, 2016, 8, 107.	4.1	74
10	Content and Performance of the MiniMUGA Genotyping Array: A New Tool To Improve Rigor and Reproducibility in Mouse Research. Genetics, 2020, 216, 905-930.	2.9	58
11	Using Animal Models to Disentangle the Role of Genetic, Epigenetic, and Environmental Influences on Behavioral Outcomes Associated with Maternal Anxiety and Depression. Frontiers in Psychiatry, 2011, 2, 44.	2.6	40
12	Gut feelings: A role for the intestinal microbiota in anorexia nervosa?. International Journal of Eating Disorders, 2015, 48, 449-451.	4.0	38
13	Prototypical anxiolytics do not reduce anxiety-like behavior in the open field in C57BL/6J mice. Pharmacology Biochemistry and Behavior, 2015, 133, 7-17.	2.9	37
14	Sequence variant analysis reveals poor correlations in microbial taxonomic abundance between humans and mice after gnotobiotic transfer. ISME Journal, 2020, 14, 1809-1820.	9.8	30
15	Sex-exclusive quantitative trait loci influences in alcohol-related phenotypes. , 1999, 88, 647-652.		29
16	Inbred Strain Variant Database (ISVdb): A Repository for Probabilistically Informed Sequence Differences Among the Collaborative Cross Strains and Their Founders. G3: Genes, Genomes, Genetics, 2017, 7, 1623-1630.	1.8	26
17	Characterization of genetically complex Collaborative Cross mouse strains that model divergent locomotor activating and reinforcing properties of cocaine. Psychopharmacology, 2020, 237, 979-996.	3.1	25
18	Quantitative trait locus and haplotype mapping in closely related inbred strains identifies a locus for open field behavior. Mammalian Genome, 2010, 21, 231-246.	2.2	24

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#	Article	IF	CITATIONS
19	Intravenous cocaine self-administration in a panel of inbred mouse strains differing in acute locomotor sensitivity to cocaine. Psychopharmacology, 2018, 235, 1179-1189.	3.1	21
20	Ovariectomy results in inbred strain-specific increases in anxiety-like behavior in mice. Physiology and Behavior, 2016, 167, 404-412.	2.1	19
21	Epistatic quantitative trait loci for alcohol preference in mice. Behavior Genetics, 2000, 30, 431-437.	2.1	14
22	Gut microbial communities from patients with anorexia nervosa do not influence body weight in recipient germ-free mice. Gut Microbes, 2021, 13, 1-15.	9.8	14
23	Prospects for finding the mechanisms of sex differences in addiction with human and model organism genetic analysis. Genes, Brain and Behavior, 2020, 19, e12645.	2.2	13
24	Reciprocal F1 Hybrids of Two Inbred Mouse Strains Reveal Parent-of-Origin and Perinatal Diet Effects on Behavior and Expression. G3: Genes, Genomes, Genetics, 2018, 8, 3447-3468.	1.8	12
25	A multivariate model for the analysis of sibship covariance structure using marker information and multiple quantitative traits. Genetic Epidemiology, 1997, 14, 921-926.	1.3	10
26	Mean-Variance QTL Mapping Identifies Novel QTL for Circadian Activity and Exploratory Behavior in Mice. G3: Genes, Genomes, Genetics, 2018, 8, 3783-3790.	1.8	10
27	Developmental loss of effect of a Chromosome 15 QTL on alcohol acceptance. Mammalian Genome, 1998, 9, 991-994.	2.2	9
28	Bayesian modeling of skewed X inactivation in genetically diverse mice identifies a novel <i>Xce</i> allele associated with copy number changes. Genetics, 2021, 218, .	2.9	5
29	Pharmacokinetic and pharmacodynamic analyses of cocaine and its metabolites in behaviorally divergent inbred mouse strains. Genes, Brain and Behavior, 2021, 20, e12666.	2.2	3
30	Behavioral characterization of a novel Cisd2 mutant mouse. Behavioural Brain Research, 2021, 405, 113187.	2.2	3
31	A multivariate model for the analysis of sibship covariance structure using marker information and multiple quantitative traits. Genetic Epidemiology, 1997, 14, 921-926.	1.3	3
32	Introduction to mammalian genome special issue: genetics of behavior. Mammalian Genome, 2014, 25, 1-2.	2.2	1
33	Contemporary neurobehavioral genetics. Pharmacology Biochemistry and Behavior, 2014, 123, 1-2.	2.9	Ο