Nathan J Bivens

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5278705/publications.pdf

Version: 2024-02-01

26 papers 1,060 citations

16 h-index 27 g-index

28 all docs 28 docs citations

times ranked

28

1789 citing authors

#	Article	IF	Citations
1	Long-Term Effects of Developmental Exposure to Oxycodone on Gut Microbiota and Relationship to Adult Behaviors and Metabolism. MSystems, 2022, 7, .	3.8	6
2	Disruption of global hypothalamic microRNA (miR) profiles and associated behavioral changes in California mice (Peromyscus californicus) developmentally exposed to endocrine disrupting chemicals. Hormones and Behavior, 2021, 128, 104890.	2.1	17
3	Developmental exposure to silver nanoparticles leads to long term gut dysbiosis and neurobehavioral alterations. Scientific Reports, 2021, 11, 6558.	3.3	22
4	Single Nucleus RNA Sequence (snRNAseq) Analysis of the Spectrum of Trophoblast Lineages Generated From Human Pluripotent Stem Cells in vitro. Frontiers in Cell and Developmental Biology, 2021, 9, 695248.	3.7	12
5	Placental Changes in the serotonin transporter (Slc6a4) knockout mouse suggest a role for serotonin in controlling nutrient acquisition. Placenta, 2021, 115, 158-168.	1.5	8
6	miRNA changes in the mouse placenta due to bisphenol A exposure. Epigenomics, 2021, 13, 1909-1919.	2.1	8
7	Influence of PCR cycle number on 16S rRNA gene amplicon sequencing of low biomass samples. Journal of Microbiological Methods, 2020, 176, 106033.	1.6	23
8	Maternal oxycodone treatment causes pathophysiological changes in the mouse placenta. Placenta, 2020, 100, 96-110.	1.5	16
9	Changes in nucleus accumbens gene expression accompany sex-specific suppression of spontaneous physical activity in aromatase knockout mice. Hormones and Behavior, 2020, 121, 104719.	2.1	8
10	Developmental exposure of California mice to endocrine disrupting chemicals and potential effects on the microbiome-gut-brain axis at adulthood. Scientific Reports, 2020, 10, 10902.	3.3	23
11	Bisphenol A and bisphenol S disruptions of the mouse placenta and potential effects on the placenta–brain axis. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 4642-4652.	7.1	92
12	Sexual dimorphism in brain transcriptomes of Amami spiny rats (Tokudaia osimensis): a rodent species where males lack the Y chromosome. BMC Genomics, 2019, 20, 87.	2.8	4
13	Early genistein exposure of California mice and effects on the gut microbiota–brain axis. Journal of Endocrinology, 2019, 242, 139-157.	2.6	21
14	Transcriptomic alterations in the brain of painted turtles (<i>Chrysemys picta</i>) developmentally exposed to bisphenol A or ethinyl estradiol. Physiological Genomics, 2017, 49, 201-215.	2.3	18
15	Bisphenol A (BPA) in the serum of pet dogs following short-term consumption of canned dog food and potential health consequences of exposure to BPA. Science of the Total Environment, 2017, 579, 1804-1814.	8.0	43
16	Consumption of a high-fat diet alters the seminal fluid and gut microbiomes in male mice. Reproduction, Fertility and Development, 2017, 29, 1602.	0.4	38
17	Gut Dysbiosis and Neurobehavioral Alterations in Rats Exposed to Silver Nanoparticles. Scientific Reports, 2017, 7, 2822.	3.3	91
18	Hypothalamic transcriptomic alterations in male and female California mice (<i>Peromyscus) Tj ETQq0 0 0 rgBT 2017, 5, e13133.</i>	/Overlock 1.7	10 Tf 50 67 To 27

2017, 5, e13133.

#	Article	lF	CITATIONS
19	RNAâ€Seq Library Construction Methods for Transcriptome Analysis. Current Protocols in Plant Biology, 2016, 1, 197-215.	2.8	8
20	Effects of exposure to bisphenol A and ethinyl estradiol on the gut microbiota of parents and their offspring in a rodent model. Gut Microbes, 2016, 7, 471-485.	9.8	121
21	Discovery of a Novel Seminal Fluid Microbiome and Influence of Estrogen Receptor Alpha Genetic Status. Scientific Reports, 2016, 6, 23027.	3.3	59
22	Evaluation of Commercially Available RNA Amplification Kits for RNA Sequencing Using Very Low Input Amounts of Total RNA. Journal of Biomolecular Techniques, 2015, 26, 4-18.	1.5	46
23	Genome Sequences of <i>Pseudoalteromonas</i> Strains ATCC BAA-314, ATCC 70018, and ATCC 70019. Genome Announcements, 2015, 3, .	0.8	3
24	Effects of Vendor and Genetic Background on the Composition of the Fecal Microbiota of Inbred Mice. PLoS ONE, 2015, 10, e0116704.	2.5	268
25	Large-Scale Generation and Analysis of Expressed Sequence Tags from Porcine Ovary1. Biology of Reproduction, 2004, 71, 1991-2002.	2.7	19
26	Developmental Expression of 2489 Gene Clusters During Pig Embryogenesis: An Expressed Sequence Tag Project1. Biology of Reproduction, 2004, 71, 1230-1243.	2.7	53