James W Macdonald

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5902887/publications.pdf

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41 papers 4,374 citations

430874 18 h-index 302126 39 g-index

43 all docs 43 docs citations

43 times ranked

11729 citing authors

#	Article	IF	CITATIONS
1	Orchestrating high-throughput genomic analysis with Bioconductor. Nature Methods, 2015, 12, 115-121.	19.0	3,070
2	Human Organ-Specific Endothelial Cell Heterogeneity. IScience, 2018, 4, 20-35.	4.1	181
3	Genomic and transcriptomic association studies identify 16 novel susceptibility loci for venous thromboembolism. Blood, 2019, 134, 1645-1657.	1.4	162
4	Uterine overdistention induces preterm labor mediated by inflammation: observations in pregnant women and nonhuman primates. American Journal of Obstetrics and Gynecology, 2015, 213, 830.e1-830.e19.	1.3	113
5	Group B <i>Streptococcus</i> circumvents neutrophils and neutrophil extracellular traps during amniotic cavity invasion and preterm labor. Science Immunology, 2016, 1, .	11.9	84
6	Elevated CO ₂ impairs olfactoryâ€mediated neural and behavioral responses and gene expression in oceanâ€phase coho salmon (<i>Oncorhynchus kisutch</i>). Global Change Biology, 2019, 25, 963-977.	9.5	77
7	Infant sex-specific placental cadmium and DNA methylation associations. Environmental Research, 2015, 138, 74-81.	7.5	63
8	Long-term outdoor air pollution and DNA methylation in circulating monocytes: results from the Multi-Ethnic Study of Atherosclerosis (MESA). Environmental Health, $2016,15,119.$	4.0	62
9	Silymarin Suppresses Cellular Inflammation By Inducing Reparative Stress Signaling. Journal of Natural Products, 2015, 78, 1990-2000.	3.0	53
10	Histone Deacetylase Inhibitors Antagonize Distinct Pathways to Suppress Tumorigenesis of Embryonal Rhabdomyosarcoma. PLoS ONE, 2015, 10, e0144320.	2.5	51
11	The Epigenetic Factor Landscape of Developing Neocortex Is Regulated by Transcription Factors Pax6â†' Tbr2â†' Tbr1. Frontiers in Neuroscience, 2018, 12, 571.	2.8	46
12	Autophagy accounts for approximately one-third of mitochondrial protein turnover and is protein selective. Autophagy, 2019, 15, 1592-1605.	9.1	41
13	A Combination Therapy of Nicotinamide and Progesterone Improves Functional Recovery following Traumatic Brain Injury. Journal of Neurotrauma, 2015, 32, 765-779.	3.4	31
14	Genetic determinants of susceptibility to silver nanoparticleâ€induced acute lung inflammation in mice. FASEB Journal, 2017, 31, 4600-4611.	0.5	28
15	PI3K-Yap activity drives cortical gyrification and hydrocephalus in mice. ELife, 2019, 8, .	6.0	28
16	Choriodecidual Group B Streptococcal Infection Induces miR-155-5p in the Fetal Lung in Macaca nemestrina. Infection and Immunity, 2015, 83, 3909-3917.	2.2	26
17	Evidence of cardiac involvement in the fetal inflammatory response syndrome: disruption of gene networks programming cardiac development in nonhuman primates. American Journal of Obstetrics and Gynecology, 2018, 218, 438.e1-438.e16.	1.3	23
18	<i>In utero (i) exposure to diesel exhaust particulates is associated with an altered cardiac transcriptional response to transverse aortic constriction and altered DNA methylation. FASEB Journal, 2017, 31, 4935-4945.</i>	0.5	20

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19	Human Cervical Mucus Plugs Exhibit Insufficiencies in Antimicrobial Activity Towards Group B Streptococcus. Journal of Infectious Diseases, 2018, 217, 1626-1636.	4.0	19
20	$\hat{l}\pm7$ Nicotinic Acetylcholine Receptor Signaling Modulates Ovine Fetal Brain Astrocytes Transcriptome in Response to Endotoxin. Frontiers in Immunology, 2019, 10, 1063.	4.8	18
21	Spiny mice activate unique transcriptional programs after severe kidney injury regenerating organ function without fibrosis. IScience, 2021, 24, 103269.	4.1	17
22	Network analysis of microRNA and mRNA seasonal dynamics in a highly plastic sensorimotor neural circuit. BMC Genomics, 2015, 16, 905.	2.8	15
23	Transcriptional analysis of human cranial compartments with different embryonic origins. Archives of Oral Biology, 2015, 60, 1450-1460.	1.8	13
24	From the Cover: Cadmium Exposure Differentially Alters Odorant-Driven Behaviors and Expression of Olfactory Receptors in Juvenile Coho Salmon <i>(Oncorhynchus kisutch)</i>). Toxicological Sciences, 2016, 154, 267-277.	3.1	13
25	Short Report: Circulating microRNAs are associated with incident diabetes over 10 years in Japanese Americans. Scientific Reports, 2020, 10, 6509.	3.3	12
26	Multiâ€phenotype analyses of hemostatic traits with cardiovascular events reveal novel genetic associations. Journal of Thrombosis and Haemostasis, 2022, 20, 1331-1349.	3.8	12
27	Epigenome-wide analysis of long-term air pollution exposure and DNA methylation in monocytes: results from the Multi-Ethnic Study of Atherosclerosis. Epigenetics, 2022, 17, 1-17.	2.7	11
28	Neonatal Exposure to BPA, BDE-99, and PCB Produces Persistent Changes in Hepatic Transcriptome Associated With Gut Dysbiosis in Adult Mouse Livers. Toxicological Sciences, 2021, 184, 83-103.	3.1	10
29	Dried blood spot compared to plasma measurements of blood-based biomarkers of brain injury in neonatal encephalopathy. Pediatric Research, 2019, 85, 655-661.	2.3	9
30	Transcriptome data of temporal and cingulate cortex in the Rett syndrome brain. Scientific Data, 2020, 7, 192.	5.3	9
31	Persistence of improved glucose homeostasis in Gclm null mice with age and cadmium treatment. Redox Biology, 2022, 49, 102213.	9.0	9
32	Repeated Gestational Exposure of Mice to Chlorpyrifos Oxon Is Associated with Paraoxonase 1 (PON1) Modulated Effects in Maternal and Fetal Tissues. Toxicological Sciences, 2014, 141, 409-422.	3.1	8
33	Genetic associations of breast and prostate cancer are enriched for regulatory elements identified in disease-related tissues. Human Genetics, 2019, 138, 1091-1104.	3.8	7
34	Whole genome methylation and transcriptome analyses to identify risk for cerebral palsy (CP) in extremely low gestational age neonates (ELGAN). Scientific Reports, 2021, 11, 5305.	3.3	7
35	Impact of processing methods on urinary biomarkers analysis in neonates. Pediatric Nephrology, 2018, 33, 181-186.	1.7	5
36	Paraoxonase 2 deficiency in mice alters motor behavior and causes region-specific transcript changes in the brain. Neurotoxicology and Teratology, 2021, 87, 107010.	2.4	5

#	Article	lF	CITATIONS
37	Extracellular vesicle microRNAs as predictors of response to omalizumab in chronic spontaneous urticaria. Allergy: European Journal of Allergy and Clinical Immunology, 2021, 76, 1274-1277.	5.7	4
38	Associations of plasma miRNAs with waist circumference and insulin resistance among women with polycystic ovary syndrome – Pilot study. Molecular and Cellular Endocrinology, 2022, 554, 111723.	3.2	4
39	Temporal changes in the brain lipidome during neurodevelopment of Smith–Lemli–Opitz syndrome mice. Analyst, The, 2022, , .	3.5	2
40	Unique basophil microRNA signature in chronic spontaneous urticaria patients who respond to omalizumab. Clinical and Experimental Allergy, 2021, 51, 1648-1652.	2.9	1
41	Associations of prenatal exposure to NO2 and near roadway residence with placental gene expression. ISEE Conference Abstracts, 2021, 2021, .	0.0	0