Richard S Lee

List of Publications by Year in descending order

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Version: 2024-02-01

394421 315739 1,573 39 19 38 citations h-index g-index papers 39 39 39 2630 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Prenatal stress enhances atherosclerosis and telomere shortening in ApoE knockout mouse offspring. American Journal of Physiology - Regulatory Integrative and Comparative Physiology, 2022, 323, R68-R80.	1.8	1
2	Characterization of glucocorticoid-induced loss of DNA methylation of the stress-response gene Fkbp5 in neuronal cells. Epigenetics, 2021, 16, 1-21.	2.7	5
3	Methylomic and transcriptomic predictors of one-month exposure to cortisol in healthy individuals. Stress, 2021, 24, 840-848.	1.8	0
4	Cross-species Association Between Telomere Length and Glucocorticoid Exposure. Journal of Clinical Endocrinology and Metabolism, 2021, 106, e5124-e5135.	3. 6	6
5	DNA methylation and behavioral dysfunction in males with 47,XXY and 49,XXXXY: a pilot study. Clinical Epigenetics, 2021, 13, 136.	4.1	2
6	Fluoxetine and environmental enrichment similarly reverse chronic social stress-related depressionand anxiety-like behavior, but have differential effects on amygdala gene expression. Neurobiology of Stress, 2021, 15, 100392.	4.0	17
7	Genetically-predicted prefrontal DRD4 gene expression modulates differentiated brain responses to food cues in adolescent girls and boys. Scientific Reports, 2021, 11, 24094.	3.3	5
8	Neuropeptides and oligopeptidases in schizophrenia. Neuroscience and Biobehavioral Reviews, 2020, 108, 679-693.	6.1	24
9	IncRNAKB, a knowledgebase of tissue-specific functional annotation and trait association of long noncoding RNA. Scientific Data, 2020, 7, 326.	5. 3	40
10	Evaluation of NDEL1 oligopeptidase activity in blood and brain in an animal model of schizophrenia: effects of psychostimulants and antipsychotics. Scientific Reports, 2020, 10, 18513.	3.3	6
11	Tubulin Polymerization Promoting Protein (TPPP) gene methylation and corpus callosum measures in maltreated children. Psychiatry Research - Neuroimaging, 2020, 298, 111058.	1.8	4
12	Predicted DRD4 prefrontal gene expression moderates snack intake and stress perception in response to the environment in adolescents. PLoS ONE, 2020, 15, e0234601.	2.5	9
13	Prenatal stress enhances NNK-induced lung tumors in A/J mice. Carcinogenesis, 2020, 41, 1713-1723.	2.8	4
14	Differentially methylated regions in bipolar disorder and suicide. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2019, 180, 496-507.	1.7	19
15	Disseminated sporotrichosis following iatrogenic immunosuppression for suspected pyoderma gangrenosum. Lancet Infectious Diseases, The, 2019, 19, e385-e391.	9.1	32
16	Extracellular Matrix Protein Tenascin C Increases Phagocytosis Mediated by CD47 Loss of Function in Glioblastoma. Cancer Research, 2019, 79, 2697-2708.	0.9	48
17	Effect of Genotype and Maternal Affective Disorder on Intronic Methylation of FK506 Binding Protein 5 in Cord Blood DNA. Frontiers in Genetics, 2018, 9, 648.	2.3	9
18	A Rat Methyl-Seq Platform to Identify Epigenetic Changes Associated with Stress Exposure. Journal of Visualized Experiments, 2018, , .	0.3	3

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19	DNA methylation and sex-specific expression of FKBP5 as correlates of one-month bedtime cortisol levels in healthy individuals. Psychoneuroendocrinology, 2018, 97, 164-173.	2.7	30
20	Type 2 diabetes and cardiometabolic risk may be associated with increase in DNA methylation of FKBP5. Clinical Epigenetics, 2018, 10, 82.	4.1	23
21	Methylation in OTX2 and related genes, maltreatment, and depression in children. Neuropsychopharmacology, 2018, 43, 2204-2211.	5.4	38
22	Early Life Stress as a Predictor of Co-Occurring Alcohol Use Disorder and Post-Traumatic Stress Disorder. Alcohol Research: Current Reviews, 2018, 39, 147-159.	3.6	10
23	Genome-wide Methyl-Seq analysis of blood-brain targets of glucocorticoid exposure. Epigenetics, 2017, 12, 637-652.	2.7	39
24	Saccharomyces boulardii probiotic-associated fungemia: questioning the safety of this preventive probiotic's use. Diagnostic Microbiology and Infectious Disease, 2017, 87, 286-288.	1.8	37
25	TET1 deficiency attenuates the DNA damage response and promotes resistance to DNA damaging agents. Epigenetics, 2017, 12, 854-864.	2.7	20
26	Can Multidrug-Resistant Candidaauris Be Reliably Identified in Clinical Microbiology Laboratories?. Journal of Clinical Microbiology, 2017, 55, 638-640.	3.9	181
27	Reduced DNA methylation of FKBP5 in Cushing's syndrome. Endocrine, 2016, 54, 768-777.	2.3	37
28	A critical period of vulnerability to adolescent stress: epigenetic mediators in mesocortical dopaminergic neurons. Human Molecular Genetics, 2016, 25, 1370-1381.	2.9	41
29	Failure to upregulate Agrp and Orexin in response to activity based anorexia in weight loss vulnerable rats characterized by passive stress coping and prenatal stress experience. Psychoneuroendocrinology, 2016, 67, 171-181.	2.7	24
30	Adaptation of the targeted capture Methyl-Seq platform for the mouse genome identifies novel tissue-specific DNA methylation patterns of genes involved in neurodevelopment. Epigenetics, 2015, 10, 581-596.	2.7	18
31	Alterations in DNA methylation of Fkbp5 as a determinant of blood–brain correlation of glucocorticoid exposure. Psychoneuroendocrinology, 2014, 44, 112-122.	2.7	101
32	Prenatal stress decreases <i>Bdnf</i> expression and increases methylation of <i>Bdnf</i> exon IV in rats. Epigenetics, 2014, 9, 437-447.	2.7	154
33	Questionable Utility of Galactomannan Testing for Diagnosis of Exserohilum rostratum Infection. Journal of Clinical Microbiology, 2014, 52, 2742-2743.	3.9	2
34	Environmental Stressors and Epigenetic Control of the Hypothalamic-Pituitary-Adrenal Axis. Neuroendocrinology, 2014, 100, 278-287.	2.5	54
35	Proper Surgical Training and Grading of Complications for Shang Ring Circumcision Are Necessary. Journal of Acquired Immune Deficiency Syndromes (1999), 2013, 64, e11.	2.1	7
36	A measure of glucocorticoid load provided by DNA methylation of Fkbp5 in mice. Psychopharmacology, 2011, 218, 303-312.	3.1	100

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37	Adaptation of the CHARM DNA methylation platform for the rat genome reveals novel brain region-specific differences. Epigenetics, 2011, 6, 1378-1390.	2.7	17
38	Chronic Corticosterone Exposure Increases Expression and Decreases Deoxyribonucleic Acid Methylation of Fkbp5 in Mice. Endocrinology, 2010, 151, 4332-4343.	2.8	248
39	A Genome-Wide Screen for Normally Methylated Human CpG Islands That Can Identify Novel Imprinted Genes. Genome Research, 2002, 12, 543-554.	5 . 5	158