

Joseph M Dhahbi

List of Publications by Year in descending order

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Version: 2024-02-01

38
papers

2,149
citations

236925

25
h-index

330143

37
g-index

39
all docs

39
docs citations

39
times ranked

3126
citing authors

#	ARTICLE	IF	CITATIONS
1	Plasma miRNA Profile of Crohn's Disease and Rheumatoid Arthritis Patients. <i>Biology</i> , 2022, 11, 508.	2.8	2
2	Specific PIWI-Interacting RNAs and Related Small Noncoding RNAs Are Associated With Ovarian Aging in Ames Dwarf (df/df) Mice. <i>Journals of Gerontology - Series A Biological Sciences and Medical Sciences</i> , 2021, 76, 1561-1570.	3.6	3
3	Organ reserve, excess metabolic capacity, and aging. <i>Biogerontology</i> , 2018, 19, 171-184.	3.9	32
4	Data Mining of Small RNA-Seq Suggests an Association Between Prostate Cancer and Altered Abundance of 5â€² Transfer RNA Halves in Seminal Fluid and Prostatic Tissues. <i>Biomarkers in Cancer</i> , 2018, 10, 1179299X1875954.	3.6	10
5	Caloric restriction impacts plasma microRNA's in rhesus monkeys. <i>Aging Cell</i> , 2017, 16, 1200-1203.	6.7	27
6	MicroRNAs Circulate in the Hemolymph of <i>Drosophila</i> and Accumulate Relative to Tissue microRNAs in an Age-Dependent Manner. <i>Genomics Insights</i> , 2016, 9, GEI.S38147.	3.0	17
7	Small Noncoding RNAs in Senescence and Aging. <i>Healthy Ageing and Longevity</i> , 2016, , 287-312.	0.2	1
8	Circulating microRNA signature of genotypeâ€œbyâ€œage interactions in the long-lived <i>A. mes dwarf</i> mouse. <i>Aging Cell</i> , 2015, 14, 1055-1066.	6.7	54
9	Circulating small non coding RNA signature in head and neck squamous cell carcinoma. <i>Oncotarget</i> , 2015, 6, 19246-19263.	1.8	89
10	Combined activation of the energy and cellular-defense pathways may explain the potent anti-senescence activity of methylene blue. <i>Redox Biology</i> , 2015, 6, 426-435.	9.0	28
11	Nordihydroguaiaretic Acid Extends the Lifespan of <i>Drosophila</i> and Mice, Increases Mortality-Related Tumors and Hemorrhagic Diathesis, and Alters Energy Homeostasis in Mice. <i>Journals of Gerontology - Series A Biological Sciences and Medical Sciences</i> , 2015, 70, 1479-1489.	3.6	19
12	Differential Effects of Hepatocyte Nuclear Factor 4Î± Isoforms on Tumor Growth and T-Cell Factor 4/AP-1 Interactions in Human Colorectal Cancer Cells. <i>Molecular and Cellular Biology</i> , 2015, 35, 3471-3490.	2.3	57
13	Circulating small noncoding RNAs as biomarkers of aging. <i>Ageing Research Reviews</i> , 2014, 17, 86-98.	10.9	74
14	Phosphorylation of p53 by TAF1 Inactivates p53-Dependent Transcription in the DNA Damage Response. <i>Molecular Cell</i> , 2014, 53, 63-74.	9.7	46
15	Deep Sequencing of Serum Small RNAs Identifies Patterns of 5â€² tRNA Half and YRNA Fragment Expression Associated with Breast Cancer. <i>Biomarkers in Cancer</i> , 2014, 6, BIC.S20764.	3.6	144
16	5â€² tRNA halves are present as abundant complexes in serum, concentrated in blood cells, and modulated by aging and calorie restriction. <i>BMC Genomics</i> , 2013, 14, 298.	2.8	204
17	Î²1-Adrenergic receptor blockade extends the life span of <i>Drosophila</i> and long-lived mice. <i>Age</i> , 2013, 35, 2099-2109.	3.0	34
18	5â€²-YRNA fragments derived by processing of transcripts from specific YRNA genes and pseudogenes are abundant in human serum and plasma. <i>Physiological Genomics</i> , 2013, 45, 990-998.	2.3	98

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19	Deep sequencing identifies circulating mouse miRNAs that are functionally implicated in manifestations of aging and responsive to calorie restriction. <i>Aging</i> , 2013, 5, 130-141.	3.1	67
20	mRNA-Seq reveals complex patterns of gene regulation and expression in the mouse skeletal muscle transcriptome associated with calorie restriction. <i>Physiological Genomics</i> , 2012, 44, 331-344.	2.3	15
21	Novel Protein Kinase Signaling Systems Regulating Lifespan Identified by Small Molecule Library Screening Using <i>Drosophila</i> . <i>PLoS ONE</i> , 2012, 7, e29782.	2.5	26
22	Statin Treatment Increases Lifespan and Improves Cardiac Health in <i>Drosophila</i> by Decreasing Specific Protein Prenylation. <i>PLoS ONE</i> , 2012, 7, e39581.	2.5	54
23	Deep Sequencing Reveals Novel MicroRNAs and Regulation of MicroRNA Expression during Cell Senescence. <i>PLoS ONE</i> , 2011, 6, e20509.	2.5	73
24	Conserved and Tissue-Specific Genic and Physiologic Responses to Caloric Restriction and Altered IGF1 Signaling in Mitotic and Postmitotic Tissues. <i>Annual Review of Nutrition</i> , 2007, 27, 193-217.	10.1	45
25	Gene Expression and Physiologic Responses of the Heart to the Initiation and Withdrawal of Caloric Restriction. <i>Journals of Gerontology - Series A Biological Sciences and Medical Sciences</i> , 2006, 61, 218-231.	3.6	76
26	Identification of potential caloric restriction mimetics by microarray profiling. <i>Physiological Genomics</i> , 2005, 23, 343-350.	2.3	144
27	Additive regulation of hepatic gene expression by dwarfism and caloric restriction. <i>Physiological Genomics</i> , 2004, 17, 307-315.	2.3	136
28	Temporal linkage between the phenotypic and genomic responses to caloric restriction. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2004, 101, 5524-5529.	7.1	234
29	Hepatic Gene Expression Profiling of Streptozotocin-Induced Diabetes. <i>Diabetes Technology and Therapeutics</i> , 2003, 5, 411-420.	4.4	25
30	Protein turnover, energy metabolism, aging, and caloric restriction. <i>Advances in Cell Aging and Gerontology</i> , 2003, , 69-86.	0.1	9
31	Aging of the Liver. , 2003, , 271-291.		5
32	Postprandial Induction of Chaperone Gene Expression Is Rapid in Mice. <i>Journal of Nutrition</i> , 2002, 132, 31-37.	2.9	17
33	Chaperone-Mediated Regulation of Hepatic Protein Secretion by Caloric Restriction. <i>Biochemical and Biophysical Research Communications</i> , 2001, 284, 335-339.	2.1	19
34	Caloric restriction alters the feeding response of key metabolic enzyme genes. <i>Mechanisms of Ageing and Development</i> , 2001, 122, 1033-1048.	4.6	97
35	Calories and aging alter gene expression for gluconeogenic, glycolytic, and nitrogen-metabolizing enzymes. <i>American Journal of Physiology - Endocrinology and Metabolism</i> , 1999, 277, E352-E360.	3.5	64
36	Dietary Energy Tissue-Specifically Regulates Endoplasmic Reticulum Chaperone Gene Expression in the Liver of Mice. , <i>Journal of Nutrition</i> , 1997, 127, 1758-1764.	2.9	49

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37	Dietary Energy Restriction in Mice Negatively Regulates Hepatic Glucose-Regulated Protein 78 (GRP78) Expression at the Posttranscriptional Level. <i>Journal of Nutrition</i> , 1996, 126, 416-423.	2.9	22
38	Dietary Calorie Restriction in Mice Induces Carbamyl Phosphate Synthetase I Gene Transcription Tissue Specifically. <i>Journal of Biological Chemistry</i> , 1996, 271, 3500-3506.	3.4	33