

Vadim E Fraifeld

List of Publications by Year in descending order

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Version: 2024-02-01

69
papers

3,538
citations

201575

27
h-index

149623

56
g-index

77
all docs

77
docs citations

77
times ranked

4801
citing authors

#	ARTICLE	IF	CITATIONS
1	Transplantation of mesenchymal stem cells causes long-term alleviation of schizophrenia-like behaviour coupled with increased neurogenesis. <i>Molecular Psychiatry</i> , 2021, 26, 4448-4463.	4.1	9
2	Background radiation impacts human longevity and cancer mortality: reconsidering the linear no-threshold paradigm. <i>Biogerontology</i> , 2021, 22, 189-195.	2.0	7
3	Machine Learning Analysis of Longevity-Associated Gene Expression Landscapes in Mammals. <i>International Journal of Molecular Sciences</i> , 2021, 22, 1073.	1.8	6
4	Expression Profiling Suggests Loss of Surface Integrity and Failure of Regenerative Repair as Major Driving Forces for Chronic Obstructive Pulmonary Disease Progression. <i>American Journal of Respiratory Cell and Molecular Biology</i> , 2021, 64, 441-452.	1.4	9
5	Systems biology analysis of lung fibrosis-related genes in the bleomycin mouse model. <i>Scientific Reports</i> , 2021, 11, 19269.	1.6	7
6	Hypercapnia-inducible factor: a hypothesis. <i>Ageing & Longevity</i> , 2021, 2, 27-31.	0.1	1
7	Small molecules for cell reprogramming: a systems biology analysis. <i>Aging</i> , 2021, 13, 25739-25762.	1.4	8
8	Metabolic remodelling of mice by hypoxic-hypercapnic environment: imitating the naked mole-rat. <i>Biogerontology</i> , 2020, 21, 143-153.	2.0	12
9	SynergyAge, a curated database for synergistic and antagonistic interactions of longevity-associated genes. <i>Scientific Data</i> , 2020, 7, 366.	2.4	16
10	Gray whale transcriptome reveals longevity adaptations associated with DNA repair and ubiquitination. <i>Aging Cell</i> , 2020, 19, e13158.	3.0	27
11	A multidimensional systems biology analysis of cellular senescence in aging and disease. <i>Genome Biology</i> , 2020, 21, 91.	3.8	177
12	c-Met as a new marker of cellular senescence. <i>Aging</i> , 2019, 11, 2889-2897.	1.4	11
13	iPSCs-Induced Cellular Reprogramming. , 2019, , .		1
14	Human Ageing Genomic Resources: new and updated databases. <i>Nucleic Acids Research</i> , 2018, 46, D1083-D1090.	6.5	511
15	The role of cellular senescence in aging through the prism of Koch-like criteria. <i>Ageing Research Reviews</i> , 2018, 41, 18-33.	5.0	36
16	Middle age enhances expression of innate immunity genes in a female mouse model of pulmonary fibrosis. <i>Biogerontology</i> , 2017, 18, 253-262.	2.0	6
17	Differential decrease in soluble and DNA-bound telomerase in senescent human fibroblasts. <i>Biogerontology</i> , 2017, 18, 525-533.	2.0	8
18	The DrugAge database of aging-related drugs. <i>Aging Cell</i> , 2017, 16, 594-597.	3.0	121

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19	Wide-scale comparative analysis of longevity genes and interventions. <i>Aging Cell</i> , 2017, 16, 1267-1275.	3.0	39
20	De novo assembling and primary analysis of genome and transcriptome of gray whale <i>Eschrichtius robustus</i> . <i>BMC Evolutionary Biology</i> , 2017, 17, 258.	3.2	11
21	A review of the biomedical innovations for healthy longevity. <i>Aging</i> , 2017, 9, 7-25.	1.4	18
22	Transplantation of mesenchymal stem cells reverses behavioural deficits and impaired neurogenesis caused by prenatal exposure to valproic acid. <i>Oncotarget</i> , 2017, 8, 17443-17452.	0.8	35
23	Tissue repair genes: the TiRe database and its implication for skin wound healing. <i>Oncotarget</i> , 2016, 7, 21145-21155.	0.8	20
24	Middle age has a significant impact on gene expression during skin wound healing in male mice. <i>Biogerontology</i> , 2016, 17, 763-770.	2.0	4
25	MitoAge: a database for comparative analysis of mitochondrial DNA, with a special focus on animal longevity. <i>Nucleic Acids Research</i> , 2016, 44, D1262-D1265.	6.5	25
26	Wound healing and longevity: Lessons from long-lived $\hat{\pm}$ MUPA mice. <i>Aging</i> , 2015, 7, 167-176.	1.4	14
27	Preferential anti-proliferative activity of <i>Varthemia iphionoides</i> (<i>Chiliadenus iphionoides</i>). <i>Israel Journal of Plant Sciences</i> , 2015, 62, 229-233.	0.3	7
28	Cellular Senescence Markers p16INK4a and p21CIP1/WAF Are Predictors of Hodgkin Lymphoma Outcome. <i>Clinical Cancer Research</i> , 2015, 21, 5164-5172.	3.2	33
29	ShcC proteins: Brain aging and beyond. <i>Ageing Research Reviews</i> , 2015, 19, 34-42.	5.0	7
30	Geroprotectors.org: a new, structured and curated database of current therapeutic interventions in aging and age-related disease. <i>Aging</i> , 2015, 7, 616-628.	1.4	93
31	Cellular senescence-like features of lung fibroblasts derived from idiopathic pulmonary fibrosis patients. <i>Aging</i> , 2015, 7, 664-672.	1.4	132
32	Age-related diseases: common or diverse pathways?. <i>Biogerontology</i> , 2014, 15, 543-545.	2.0	4
33	Uncovering the Geroprotective Potential of Medicinal Plants from the Judea Region of Israel. <i>Rejuvenation Research</i> , 2014, 17, 134-139.	0.9	11
34	LongevityMap: a database of human genetic variants associated with longevity. <i>Trends in Genetics</i> , 2013, 29, 559-560.	2.9	92
35	The role of DNA damage and repair in aging through the prism of Koch-like criteria. <i>Ageing Research Reviews</i> , 2013, 12, 661-684.	5.0	290
36	Telomere length and body temperature-independent determinants of mammalian longevity?. <i>Frontiers in Genetics</i> , 2013, 4, 111.	1.1	19

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37	Human Ageing Genomic Resources: Integrated databases and tools for the biology and genetics of ageing. <i>Nucleic Acids Research</i> , 2012, 41, D1027-D1033.	6.5	467
38	Co-regulation of polar mRNA transport and lifespan in budding yeast <i>Saccharomyces cerevisiae</i> . <i>Cell Cycle</i> , 2012, 11, 4275-4280.	1.3	3
39	Gadd45 proteins: Relevance to aging, longevity and age-related pathologies. <i>Ageing Research Reviews</i> , 2012, 11, 51-66.	5.0	126
40	Prediction of <i>C. elegans</i> Longevity Genes by Human and Worm Longevity Networks. <i>PLoS ONE</i> , 2012, 7, e48282.	1.1	49
41	In Memory of Amir Abramovich. <i>Rejuvenation Research</i> , 2011, 14, 105-106.	0.9	0
42	Linking cell polarity, aging and rejuvenation. <i>Biogerontology</i> , 2011, 12, 167-175.	2.0	17
43	Is rate of skin wound healing associated with aging or longevity phenotype?. <i>Biogerontology</i> , 2011, 12, 591-597.	2.0	24
44	Molecular links between cellular senescence, longevity and age-related diseases – a systems biology perspective. <i>Aging</i> , 2011, 3, 1178-1191.	1.4	119
45	The NetAge database: a compendium of networks for longevity, age-related diseases and associated processes. <i>Biogerontology</i> , 2010, 11, 513-522.	2.0	71
46	MicroRNA-Regulated Protein-Protein Interaction Networks: How Could They Help in Searching for Pro-Longevity Targets?. <i>Rejuvenation Research</i> , 2010, 13, 373-377.	0.9	30
47	NUMT (‘New Mighty’) Hypothesis of Longevity. <i>Rejuvenation Research</i> , 2010, 13, 152-155.	0.9	8
48	Common gene signature of cancer and longevity. <i>Mechanisms of Ageing and Development</i> , 2009, 130, 33-39.	2.2	52
49	The Human Ageing Genomic Resources: online databases and tools for biogerontologists. <i>Aging Cell</i> , 2009, 8, 65-72.	3.0	173
50	The signaling hubs at the crossroad of longevity and age-related disease networks. <i>International Journal of Biochemistry and Cell Biology</i> , 2009, 41, 516-520.	1.2	91
51	Senescing Cells Share Common Features with Dedifferentiating Cells. <i>Rejuvenation Research</i> , 2009, 12, 435-443.	0.9	35
52	Do Mitochondrial DNA and Metabolic Rate Complement Each Other in Determination of the Mammalian Maximum Longevity?. <i>Rejuvenation Research</i> , 2008, 11, 409-417.	0.9	37
53	Have We Reached the Point for In Vivo Rejuvenation?. <i>Rejuvenation Research</i> , 2008, 11, 489-492.	0.9	13
54	Longevity network: Construction and implications. <i>Mechanisms of Ageing and Development</i> , 2007, 128, 117-124.	2.2	84

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55	From Disease-Oriented to Aging/Longevity-Oriented Studies. <i>Rejuvenation Research</i> , 2006, 9, 207-210.	0.9	25
56	Mitochondrial Genome Anatomy and Species-Specific Lifespan. <i>Rejuvenation Research</i> , 2006, 9, 223-226.	0.9	16
57	p66ShcA and ageing: modulation by longevity-promoting agent aurintricarboxylic acid. <i>Mechanisms of Ageing and Development</i> , 2005, 126, 249-254.	2.2	17
58	Glutathione S-transferase hGSTM3 and ageing-associated neurodegeneration: relationship to Alzheimer's disease. <i>Mechanisms of Ageing and Development</i> , 2005, 126, 309-315.	2.2	45
59	The role of Hsp90 in cell response to hyperthermia. <i>Journal of Thermal Biology</i> , 2004, 29, 509-514.	1.1	16
60	Non-prostaglandin eicosanoids in fever and anapyrexia. <i>Frontiers in Bioscience - Landmark</i> , 2004, 9, 3339.	3.0	29
61	Superoxide dismutase, catalase and glutathione peroxidase activities in the liver of young and old mice: linear regression and correlation. <i>Archives of Gerontology and Geriatrics</i> , 2002, 35, 205-214.	1.4	23
62	Evidence supporting involvement of leukotrienes in LPS-induced hypothermia in mice. <i>American Journal of Physiology - Regulatory Integrative and Comparative Physiology</i> , 1999, 276, R52-R58.	0.9	21
63	Nimesulide prevents lipopolysaccharide-induced elevation in plasma tumor necrosis factor- α in rats. <i>Life Sciences</i> , 1998, 63, PL323-PL327.	2.0	15
64	Chapter 9 Brain eicosanoids and LPS fever: species and age differences. <i>Progress in Brain Research</i> , 1998, 115, 141-157.	0.9	43
65	Tolerance to lipopolysaccharide is not related to the ability of the hypothalamus to produce prostaglandin E2. <i>Life Sciences</i> , 1997, 61, 813-818.	2.0	10
66	Dietary restriction modifies fever response in aging rats. <i>Archives of Gerontology and Geriatrics</i> , 1997, 24, 133-140.	1.4	2
67	Delayed febrile response in old rats is not associated with an inability of hypothalamus to produce prostaglandin E2. <i>Mechanisms of Ageing and Development</i> , 1995, 79, 137-140.	2.2	9
68	Is hypothalamic prostaglandin E2 involved in avian fever?. <i>Life Sciences</i> , 1995, 56, 1343-1346.	2.0	28
69	Correlative links between natural radiation and life expectancy in the US population. <i>Biogerontology</i> , 0, , .	2.0	1