Joao Pedro de Magalhaes

List of Publications by Citations

 $\textbf{Source:} \ https://exaly.com/author-pdf/5440995/joao-pedro-de-magalhaes-publications-by-citations.pdf$

Version: 2024-04-10

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

158 papers

7,845 citations

50 h-index

85 g-index

175 ext. papers

10,404 ext. citations

avg, IF

6.62 L-index

#	Paper	IF	Citations
158	Meta-analysis of age-related gene expression profiles identifies common signatures of aging. <i>Bioinformatics</i> , 2009 , 25, 875-81	7.2	496
157	Human Ageing Genomic Resources: integrated databases and tools for the biology and genetics of ageing. <i>Nucleic Acids Research</i> , 2013 , 41, D1027-33	20.1	389
156	Gene co-expression analysis for functional classification and gene-disease predictions. <i>Briefings in Bioinformatics</i> , 2018 , 19, 575-592	13.4	377
155	A database of vertebrate longevity records and their relation to other life-history traits. <i>Journal of Evolutionary Biology</i> , 2009 , 22, 1770-4	2.3	325
154	Subcytotoxic H2O2 stress triggers a release of transforming growth factor-beta 1, which induces biomarkers of cellular senescence of human diploid fibroblasts. <i>Journal of Biological Chemistry</i> , 2001 , 276, 2531-7	5.4	247
153	RNA-Seq Signatures Normalized by mRNA Abundance Allow Absolute Deconvolution of Human Immune Cell Types. <i>Cell Reports</i> , 2019 , 26, 1627-1640.e7	10.6	238
152	Human Ageing Genomic Resources: new and updated databases. <i>Nucleic Acids Research</i> , 2018 , 46, D108	3 3 ≥Ð.10	90 28
151	An analysis of the relationship between metabolism, developmental schedules, and longevity using phylogenetic independent contrasts. <i>Journals of Gerontology - Series A Biological Sciences and Medical Sciences</i> , 2007 , 62, 149-60	6.4	227
150	The role of DNA methylation in aging, rejuvenation, and age-related disease. <i>Rejuvenation Research</i> , 2012 , 15, 483-94	2.6	214
149	Insights into the evolution of longevity from the bowhead whale genome. Cell Reports, 2015, 10, 112-2	210.6	203
148	How ageing processes influence cancer. <i>Nature Reviews Cancer</i> , 2013 , 13, 357-65	31.3	199
147	Longer lifespan in male mice treated with a weakly estrogenic agonist, an antioxidant, an Eglucosidase inhibitor or a Nrf2-inducer. <i>Aging Cell</i> , 2016 , 15, 872-84	9.9	176
146	Stress, cell senescence and organismal ageing. <i>Mechanisms of Ageing and Development</i> , 2018 , 170, 2-9	5.6	152
145	The Human Ageing Genomic Resources: online databases and tools for biogerontologists. <i>Aging Cell</i> , 2009 , 8, 65-72	9.9	152
144	A review and appraisal of the DNA damage theory of ageing. <i>Mutation Research - Reviews in Mutation Research</i> , 2011 , 728, 12-22	7	141
143	HAGR: the Human Ageing Genomic Resources. <i>Nucleic Acids Research</i> , 2005 , 33, D537-43	20.1	109
142	WHISTLE: a high-accuracy map of the human N6-methyladenosine (m6A) epitranscriptome predicted using a machine learning approach. <i>Nucleic Acids Research</i> , 2019 , 47, e41	20.1	106

(2004-2012)

141	Genome-environment interactions that modulate aging: powerful targets for drug discovery. <i>Pharmacological Reviews</i> , 2012 , 64, 88-101	22.5	92	
140	flowAI: automatic and interactive anomaly discerning tools for flow cytometry data. <i>Bioinformatics</i> , 2016 , 32, 2473-80	7.2	85	
139	Whole transcriptome sequencing of the aging rat brain reveals dynamic RNA changes in the dark matter of the genome. <i>Age</i> , 2013 , 35, 763-76		82	
138	Cells discover fire: employing reactive oxygen species in development and consequences for aging. <i>Experimental Gerontology</i> , 2006 , 41, 1-10	4.5	82	
137	From cells to ageing: a review of models and mechanisms of cellular senescence and their impact on human ageing. <i>Experimental Cell Research</i> , 2004 , 300, 1-10	4.2	82	
136	GenAge: a genomic and proteomic network map of human ageing. FEBS Letters, 2004, 571, 243-7	3.8	81	
135	The Business of Anti-Aging Science. <i>Trends in Biotechnology</i> , 2017 , 35, 1062-1073	15.1	80	
134	The influence of genes on the aging process of mice: a statistical assessment of the genetics of aging. <i>Genetics</i> , 2005 , 169, 265-74	4	80	
133	GeneFriends: a human RNA-seq-based gene and transcript co-expression database. <i>Nucleic Acids Research</i> , 2015 , 43, D1124-32	20.1	77	
132	RNA sequencing reveals differential expression of mitochondrial and oxidation reduction genes in the long-lived naked mole-rat when compared to mice. <i>PLoS ONE</i> , 2011 , 6, e26729	3.7	77	
131	Next-generation sequencing in aging research: emerging applications, problems, pitfalls and possible solutions. <i>Ageing Research Reviews</i> , 2010 , 9, 315-23	12	77	
130	Programmatic features of aging originating in development: aging mechanisms beyond molecular damage?. <i>FASEB Journal</i> , 2012 , 26, 4821-6	0.9	76	
129	The DrugAge database of aging-related drugs. <i>Aging Cell</i> , 2017 , 16, 594-597	9.9	71	
128	From the Hayflick mosaic to the mosaics of ageing. Role of stress-induced premature senescence in human ageing. <i>International Journal of Biochemistry and Cell Biology</i> , 2002 , 34, 1415-29	5.6	70	
127	A meta-analysis of caloric restriction gene expression profiles to infer common signatures and regulatory mechanisms. <i>Molecular BioSystems</i> , 2012 , 8, 1339-49		69	
126	UVB-induced premature senescence of human diploid skin fibroblasts. <i>International Journal of Biochemistry and Cell Biology</i> , 2002 , 34, 1331-9	5.6	68	
125	A network pharmacology approach reveals new candidate caloric restriction mimetics in C. elegans. <i>Aging Cell</i> , 2016 , 15, 256-66	9.9	66	
124	Gene expression and regulation in H2O2-induced premature senescence of human foreskin fibroblasts expressing or not telomerase. <i>Experimental Gerontology</i> , 2004 , 39, 1379-89	4.5	65	

123	Geroprotectors.org: a new, structured and curated database of current therapeutic interventions in aging and age-related disease. <i>Aging</i> , 2015 , 7, 616-28	5.6	65
122	Being cool: how body temperature influences ageing and longevity. <i>Biogerontology</i> , 2015 , 16, 383-97	4.5	64
121	Systems biology and longevity: an emerging approach to identify innovative anti-aging targets and strategies. <i>Current Pharmaceutical Design</i> , 2010 , 16, 802-13	3.3	64
120	A review of supervised machine learning applied to ageing research. <i>Biogerontology</i> , 2017 , 18, 171-188	4.5	62
119	LongevityMap: a database of human genetic variants associated with longevity. <i>Trends in Genetics</i> , 2013 , 29, 559-60	8.5	62
118	The Naked Mole Rat Genome Resource: facilitating analyses of cancer and longevity-related adaptations. <i>Bioinformatics</i> , 2014 , 30, 3558-60	7.2	62
117	Cell divisions and mammalian aging: integrative biology insights from genes that regulate longevity. <i>BioEssays</i> , 2008 , 30, 567-78	4.1	62
116	Stress-induced premature senescence in BJ and hTERT-BJ1 human foreskin fibroblasts. <i>FEBS Letters</i> , 2002 , 523, 157-62	3.8	62
115	Genomes optimize reproduction: aging as a consequence of the developmental program. <i>Physiology</i> , 2005 , 20, 252-9	9.8	61
114	A comparison of human and mouse gene co-expression networks reveals conservation and divergence at the tissue, pathway and disease levels. <i>BMC Evolutionary Biology</i> , 2015 , 15, 259	3	56
113	The Digital Ageing Atlas: integrating the diversity of age-related changes into a unified resource. <i>Nucleic Acids Research</i> , 2015 , 43, D873-8	20.1	53
112	Systematic analysis of the gerontome reveals links between aging and age-related diseases. <i>Human Molecular Genetics</i> , 2016 , 25, 4804-4818	5.6	53
111	GeneFriends: an online co-expression analysis tool to identify novel gene targets for aging and complex diseases. <i>BMC Genomics</i> , 2012 , 13, 535	4.5	52
110	m6A-Atlas: a comprehensive knowledgebase for unraveling the N6-methyladenosine (m6A) epitranscriptome. <i>Nucleic Acids Research</i> , 2021 , 49, D134-D143	20.1	52
109	From humans to hydra: patterns of cancer across the tree of life. <i>Biological Reviews</i> , 2018 , 93, 1715-173	413.5	50
108	A multidimensional systems biology analysis of cellular senescence in aging and disease. <i>Genome Biology</i> , 2020 , 21, 91	18.3	49
107	Naked mole rats can undergo developmental, oncogene-induced and DNA damage-induced cellular senescence. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018 , 115, 1801-1806	11.5	44
106	Accelerated protein evolution analysis reveals genes and pathways associated with the evolution of mammalian longevity. <i>Age</i> , 2013 , 35, 301-14		42

(2007-2014)

105	Why genes extending lifespan in model organisms have not been consistently associated with human longevity and what it means to translation research. <i>Cell Cycle</i> , 2014 , 13, 2671-3	4.7	42
104	Biological Processes Modulating Longevity across Primates: A Phylogenetic Genome-Phenome Analysis. <i>Molecular Biology and Evolution</i> , 2018 , 35, 1990-2004	8.3	42
103	Transcriptome analysis in calorie-restricted rats implicates epigenetic and post-translational mechanisms in neuroprotection and aging. <i>Genome Biology</i> , 2015 , 16, 285	18.3	41
102	A data mining approach for classifying DNA repair genes into ageing-related or non-ageing-related. <i>BMC Genomics</i> , 2011 , 12, 27	4.5	40
101	Dissecting the gene network of dietary restriction to identify evolutionarily conserved pathways and new functional genes. <i>PLoS Genetics</i> , 2012 , 8, e1002834	6	40
100	Mitochondrially encoded methionine is inversely related to longevity in mammals. <i>Aging Cell</i> , 2011 , 10, 198-207	9.9	39
99	Open-minded scepticism: inferring the causal mechanisms of human ageing from genetic perturbations. <i>Ageing Research Reviews</i> , 2005 , 4, 1-22	12	39
98	Vive la radiorEistance!: converging research in radiobiology and biogerontology to enhance human radioresistance for deep space exploration and colonization. <i>Oncotarget</i> , 2018 , 9, 14692-14722	3.3	38
97	Cognitive aging as an extension of brain development: a model linking learning, brain plasticity, and neurodegeneration. <i>Mechanisms of Ageing and Development</i> , 2005 , 126, 1026-33	5.6	35
96	Growth kinetics rather than stress accelerate telomere shortening in cultures of human diploid fibroblasts in oxidative stress-induced premature senescence. <i>FEBS Letters</i> , 2001 , 502, 109-12	3.8	34
95	Prediction of C. elegans longevity genes by human and worm longevity networks. <i>PLoS ONE</i> , 2012 , 7, e48282	3.7	34
94	The effects of donor age on organ transplants: A review and implications for aging research. <i>Experimental Gerontology</i> , 2018 , 110, 230-240	4.5	34
93	Biohorology and biomarkers of aging: Current state-of-the-art, challenges and opportunities. <i>Ageing Research Reviews</i> , 2020 , 60, 101050	12	33
92	m7GHub: deciphering the location, regulation and pathogenesis of internal mRNA N7-methylguanosine (m7G) sites in human. <i>Bioinformatics</i> , 2020 , 36, 3528-3536	7.2	31
91	A new approach for interpreting Random Forest models and its application to the biology of ageing. <i>Bioinformatics</i> , 2018 , 34, 2449-2456	7.2	31
90	The scientific quest for lasting youth: prospects for curing aging. <i>Rejuvenation Research</i> , 2014 , 17, 458-	62 .6	31
89	RMDisease: a database of genetic variants that affect RNA modifications, with implications for epitranscriptome pathogenesis. <i>Nucleic Acids Research</i> , 2021 , 49, D1396-D1404	20.1	31
88	Analyses of human-chimpanzee orthologous gene pairs to explore evolutionary hypotheses of aging. <i>Mechanisms of Ageing and Development</i> , 2007 , 128, 355-64	5.6	30

87	A human tissue-specific transcriptomic analysis reveals a complex relationship between aging, cancer, and cellular senescence. <i>Aging Cell</i> , 2019 , 18, e13041	9.9	29
86	Age-related gene-specific changes of A-to-I mRNA editing in the human brain. <i>Mechanisms of Ageing and Development</i> , 2010 , 131, 445-7	5.6	29
85	miRNA-31 Improves Cognition and Abolishes Amyloid-Pathology by Targeting APP and BACE1 in an Animal Model of Alzheimer's Disease. <i>Molecular Therapy - Nucleic Acids</i> , 2020 , 19, 1219-1236	10.7	28
84	The evolution of mammalian aging. Experimental Gerontology, 2002, 37, 769-75	4.5	26
83	To help aging populations, classify organismal senescence. <i>Science</i> , 2019 , 366, 576-578	33.3	24
82	Cell resilience in species life spans: a link to inflammation?. <i>Aging Cell</i> , 2010 , 9, 519-26	9.9	23
81	Predicting the Pro-Longevity or Anti-Longevity Effect of Model Organism Genes with New Hierarchical Feature Selection Methods. <i>IEEE/ACM Transactions on Computational Biology and Bioinformatics</i> , 2015 , 12, 262-75	3	22
80	Machine learning for predicting lifespan-extending chemical compounds. <i>Aging</i> , 2017 , 9, 1721-1737	5.6	21
79	Telomeres and telomerase: a modern fountain of youth?. Rejuvenation Research, 2004, 7, 126-33	2.6	21
78	Winter is coming: the future of cryopreservation. <i>BMC Biology</i> , 2021 , 19, 56	7.3	20
77	The hoverfly and the wasp: A critique of the hallmarks of aging as a paradigm. <i>Ageing Research Reviews</i> , 2021 , 70, 101407	12	20
76	How bioinformatics can help reverse engineer human aging. <i>Ageing Research Reviews</i> , 2004 , 3, 125-41	12	19
75	Human disease-associated mitochondrial mutations fixed in nonhuman primates. <i>Journal of Molecular Evolution</i> , 2005 , 61, 491-7	3.1	19
74	A review of the biomedical innovations for healthy longevity. <i>Aging</i> , 2017 , 9, 7-25	5.6	18
73	Bioinformatics approaches for deciphering the epitranscriptome: Recent progress and emerging topics. <i>Computational and Structural Biotechnology Journal</i> , 2020 , 18, 1587-1604	6.8	18
72	Stress-induced premature senescence as alternative toxicological method for testing the long-term effects of molecules under development in the industry. <i>Biogerontology</i> , 2000 , 1, 179-83	4.5	17
71	Fish oil supplements, longevity and aging. <i>Aging</i> , 2016 , 8, 1578-82	5.6	17
70	Analysis of the FGF gene family provides insights into aquatic adaptation in cetaceans. <i>Scientific Reports</i> , 2017 , 7, 40233	4.9	16

(2018-2015)

69	Insights on cryoprotectant toxicity from gene expression profiling of endothelial cells exposed to ethylene glycol. <i>Cryobiology</i> , 2015 , 71, 405-12	2.7	16
68	Circulating MicroRNAs in Young Patients with Acute Coronary Syndrome. <i>International Journal of Molecular Sciences</i> , 2018 , 19,	6.3	14
67	The biology of ageing21-47		14
66	The genomics of ecological flexibility, large brains, and long lives in capuchin monkeys revealed with fecalFACS. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021 , 118,	11.5	14
65	Is mammalian aging genetically controlled?. <i>Biogerontology</i> , 2003 , 4, 119-20	4.5	13
64	Gene Size Matters: An Analysis of Gene Length in the Human Genome. <i>Frontiers in Genetics</i> , 2021 , 12, 559998	4.5	13
63	Mutational bias plays an important role in shaping longevity-related amino acid content in mammalian mtDNA-encoded proteins. <i>Journal of Molecular Evolution</i> , 2012 , 74, 332-41	3.1	12
62	A mathematical model of mortality dynamics across the lifespan combining heterogeneity and stochastic effects. <i>Experimental Gerontology</i> , 2013 , 48, 801-11	4.5	12
61	Ecological, biomedical and epidemiological approaches to understanding oxidative balance and ageing: what they can teach each other. <i>Functional Ecology</i> , 2010 , 24, 997-1006	5.6	12
60	Topological Characterization of Human and Mouse mC Epitranscriptome Revealed by Bisulfite Sequencing. <i>International Journal of Genomics</i> , 2018 , 2018, 1351964	2.5	11
59	A-to-I RNA editing does not change with age in the healthy male rat brain. <i>Biogerontology</i> , 2013 , 14, 39	5 -49 0	11
58	Ageing-associated changes in the expression of lncRNAs in human tissues reflect a transcriptional modulation in ageing pathways. <i>Mechanisms of Ageing and Development</i> , 2020 , 185, 111177	5.6	11
57	Has gene duplication impacted the evolution of Eutherian longevity?. Aging Cell, 2016, 15, 978-80	9.9	11
56	Gathering insights on disease etiology from gene expression profiles of healthy tissues. <i>Bioinformatics</i> , 2011 , 27, 3300-5	7.2	10
55	ConsRM: collection and large-scale prediction of the evolutionarily conserved RNA methylation sites, with implications for the functional epitranscriptome. <i>Briefings in Bioinformatics</i> , 2021 , 22,	13.4	10
54	Attention-based multi-label neural networks for integrated prediction and interpretation of twelve widely occurring RNA modifications. <i>Nature Communications</i> , 2021 , 12, 4011	17.4	10
53	Targeting immune dysfunction in aging. Ageing Research Reviews, 2021, 70, 101410	12	10
52	A Reassessment of Genes Modulating Aging in Mice Using Demographic Measurements of the Rate of Aging. <i>Genetics</i> , 2018 , 208, 1617-1630	4	9

51	Endless paces of degenerationapplying comparative genomics to study evolution's moulding of longevity. <i>EMBO Reports</i> , 2013 , 14, 661-2	6.5	9
50	Species Selection in Comparative Studies of Aging and Antiaging Research 2006 , 9-20		9
49	An integrative analysis of the age-associated multi-omic landscape across cancers. <i>Nature Communications</i> , 2021 , 12, 2345	17.4	9
48	Positive selection and gene duplications in tumour suppressor genes reveal clues about how cetaceans resist cancer. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2021 , 288, 20202592	4.4	9
47	Genome-wide patterns of genetic distances reveal candidate Loci contributing to human population-specific traits. <i>Annals of Human Genetics</i> , 2012 , 76, 142-58	2.2	8
46	No increase in senescence-associated beta-galactosidase activity in Werner syndrome fibroblasts after exposure to H2O2. <i>Annals of the New York Academy of Sciences</i> , 2004 , 1019, 375-8	6.5	8
45	Alice⊠ dilemma. <i>Futures</i> , 2004 , 36, 85-89	3.6	8
44	The fog of genetics: what is known, unknown and unknowable in the genetics of complex traits and diseases. <i>EMBO Reports</i> , 2019 , 20, e48054	6.5	7
43	Vulnerability of progeroid smooth muscle cells to biomechanical forces is mediated by MMP13. <i>Nature Communications</i> , 2020 , 11, 4110	17.4	7
42	Reductions in hypothalamic Gfap expression, glial cells and Etanycytes in lean and hypermetabolic Gnasxl-deficient mice. <i>Molecular Brain</i> , 2016 , 9, 39	4.5	6
41	Enhancing Epitranscriptome Module Detection from mA-Seq Data Using Threshold-Based Measurement Weighting Strategy. <i>BioMed Research International</i> , 2018 , 2018, 2075173	3	6
40	MYCN/LIN28B/Let-7/HMGA2 pathway implicated by meta-analysis of GWAS in suppression of post-natal proliferation thereby potentially contributing to aging. <i>Mechanisms of Ageing and Development</i> , 2013 , 134, 346-8	5.6	6
39	Paternal genome effects on aging: evidence for a role of Rasgrf1 in longevity determination?. <i>Mechanisms of Ageing and Development</i> , 2011 , 132, 72-3	5.6	6
38	A proposal to sequence genomes of unique interest for research on aging. <i>Journals of Gerontology - Series A Biological Sciences and Medical Sciences</i> , 2007 , 62, 583-4	6.4	6
37	Identification of polymorphisms in cancer patients that differentially affect survival with age. <i>Aging</i> , 2017 , 9, 2117-2136	5.6	6
36	A scan for genes associated with cancer mortality and longevity in pedigree dog breeds. Mammalian Genome, 2020 , 31, 215-227	3.2	6
35	SynergyAge, a curated database for synergistic and antagonistic interactions of longevity-associated genes. <i>Scientific Data</i> , 2020 , 7, 366	8.2	6
34	Integrative Genomics of Aging 2016 , 263-285		5

(2021-2015)

33	The big, the bad and the ugly: Extreme animals as inspiration for biomedical research. <i>EMBO Reports</i> , 2015 , 16, 771-6	6.5	5
32	An analysis and validation pipeline for large-scale RNAi-based screens. Scientific Reports, 2013, 3, 1076	4.9	5
31	A role for Ras signaling in modulating mammalian aging by the GH/IGF1 axis. Aging, 2011, 3, 336-7	5.6	5
30	Every gene can (and possibly will) be associated with cancer. Trends in Genetics, 2021,	8.5	5
29	A direct communication proposal to test the Zoo Hypothesis. Space Policy, 2016, 38, 22-26	1.4	5
28	Ageing transcriptome meta-analysis reveals similarities and differences between key mammalian tissues. <i>Aging</i> , 2021 , 13, 3313-3341	5.6	5
27	Ethical Perspectives in Biogerontology. Ethics and Health Policy, 2013, 173-188		4
26	Sex-specific aging in animals: Perspective and future directions <i>Aging Cell</i> , 2022 , e13542	9.9	4
25	The inherent challenges of classifying senescence-Response. <i>Science</i> , 2020 , 368, 595-596	33.3	3
24	Hormesis: a quest for virtuality?. Human and Experimental Toxicology, 2001 , 20, 311-4; discussion 319-20	3.4	3
23	Trans cohort metabolic reprogramming towards glutaminolysis in long-term successfully treated HIV-infection <i>Communications Biology</i> , 2022 , 5, 27	6.7	3
22	m5C-Atlas: a comprehensive database for decoding and annotating the 5-methylcytosine (m5C) epitranscriptome <i>Nucleic Acids Research</i> , 2022 , 50, D196-D203	20.1	3
21	A Multidimensional Systems Biology Analysis of Cellular Senescence in Ageing and Disease		3
20	Ageing Transcriptome Meta-Analysis Reveals Similarities Between Key Mammalian Tissues		3
19	Using deep learning to associate human genes with age-related diseases. <i>Bioinformatics</i> , 2020 , 36, 2202	2- 7 .208	3
18	Comparing enrichment analysis and machine learning for identifying gene properties that discriminate between gene classes. <i>Briefings in Bioinformatics</i> , 2020 , 21, 803-814	13.4	3
17	Evolution, structure and emerging roles of C1ORF112 in DNA replication, DNA damage responses, and cancer. <i>Cellular and Molecular Life Sciences</i> , 2021 , 78, 4365-4376	10.3	3
16	Molecular damage in aging. <i>Nature Aging</i> , 2021 , 1, 1096-1106		3

15	Human Ageing Genomic Resources: 2018 Update	2
14	Positive selection and gene duplications in tumour suppressor genes reveal clues about how cetaceans resist cancer	2
13	The evolution of ecological flexibility, large brains, and long lives: capuchin monkey genomics revealed with fecalFACS	2
12	Longevity pharmacology comes of age. <i>Drug Discovery Today</i> , 2021 , 26, 1559-1562	2
11	GeneFriends 2021: Updated co-expression databases and tools for human and mouse genes and transcripts	2
10	Osh6 links yeast vacuolar functions to lifespan extension and TOR. <i>Cell Cycle</i> , 2012 , 11, 2419 4.7	1
9	A Human Tissue-Specific Transcriptomic Analysis Reveals that Ageing Hinders Cancer and Boosts Cellular Senescence	1
8	An Integrative Analysis of the Age-Associated Genomic, Transcriptomic and Epigenetic Landscape across Cancers	1
7	The fog of genetics: Known unknowns and unknown unknowns in the genetics of complex traits and diseases	1
6	scAgeCom: a murine atlas of age-related changes in intercellular communication inferred with the package scDiffCom	1
5	Machine learning-based predictions of dietary restriction associations across ageing-related genes <i>BMC Bioinformatics</i> , 2022 , 23, 10	О
4	Histone Variant macroH2A1.1 Enhances Nonhomologous End Joining-dependent DNA Double-strand-break Repair and Reprogramming Efficiency of Human iPSCs <i>Stem Cells</i> , 2022 , 40, 35-48 ^{5.8}	O
3	A method for the permeabilization of live larvae to small molecules and cryoprotectants. <i>Fly</i> , 2020 , 14, 29-33	
2	Integrative genomics of aging 2021 , 151-171	

Ageing research in the post-genome era: new technologies for an old problem. *SEB Experimental Biology Series*, **2009**, 62, 99-115