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List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7079723/publications.pdf>

Version: 2024-02-01

20
papers

626
citations

687220

13
h-index

794469

19
g-index

31
all docs

31
docs citations

31
times ranked

1486
citing authors

#	ARTICLE	IF	CITATIONS
1	The Demographic Development of the First Farmers in Anatolia. <i>Current Biology</i> , 2016, 26, 2659-2666.	1.8	163
2	Gene expressionâ€based drug repurposing to target aging. <i>Aging Cell</i> , 2018, 17, e12819.	3.0	56
3	Common genetic associations between age-related diseases. <i>Nature Aging</i> , 2021, 1, 400-412.	5.3	55
4	Functional conservation in genes and pathways linking ageing and immunity. <i>Immunity and Ageing</i> , 2021, 18, 23.	1.8	38
5	The widespread increase in inter-individual variability of gene expression in the human brain with age. <i>Aging</i> , 2019, 11, 2253-2280.	1.4	37
6	Gene expression reversal toward pre-adult levels in the aging human brain and age-related loss of cellular identity. <i>Scientific Reports</i> , 2017, 7, 5894.	1.6	35
7	Temporal changes in the gene expression heterogeneity during brain development and aging. <i>Scientific Reports</i> , 2020, 10, 4080.	1.6	34
8	Human population dynamics and <i>Yersinia pestis</i> in ancient northeast Asia. <i>Science Advances</i> , 2021, 7, .	4.7	32
9	Using the drug-protein interactome to identify anti-ageing compounds for humans. <i>PLoS Computational Biology</i> , 2019, 15, e1006639.	1.5	30
10	Archaeogenomic analysis of the first steps of Neolithization in Anatolia and the Aegean. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2017, 284, 20172064.	1.2	26
11	Inter-tissue convergence of gene expression during ageing suggests age-related loss of tissue and cellular identity. <i>ELife</i> , 2022, 11, .	2.8	26
12	Molecular footprint of Medawarâ€™s mutation accumulation process in mammalian aging. <i>Aging Cell</i> , 2019, 18, e12965.	3.0	15
13	Identifying Potential Ageing-Modulating Drugs In Silico. <i>Trends in Endocrinology and Metabolism</i> , 2019, 30, 118-131.	3.1	15
14	Variation and Functional Impact of Neanderthal Ancestry in Western Asia. <i>Genome Biology and Evolution</i> , 2017, 9, 3516-3524.	1.1	14
15	Characterising Complex Enzyme Reaction Data. <i>PLoS ONE</i> , 2016, 11, e0147952.	1.1	13
16	Global network of computational biology communities: ISCB's Regional Student Groups breaking barriers. <i>F1000Research</i> , 2019, 8, 1574.	0.8	8
17	Transposable Element Landscape in <i>Drosophila</i> Populations Selected for Longevity. <i>Genome Biology and Evolution</i> , 2021, 13, .	1.1	6
18	Transcriptomic profiling of long- and short-lived mutant mice implicates mitochondrial metabolism in ageing and shows signatures of normal ageing in progeroid mice. <i>Mechanisms of Ageing and Development</i> , 2021, 194, 111437.	2.2	6

#	ARTICLE	IF	CITATIONS
19	Highlights from the 16th International Society for Computational Biology Student Council Symposium 2020. F1000Research, 2021, 10, 443.	0.8	4
20	Lessons from a ten-year-long journey: building a student-driven computational biology society across Turkey. F1000Research, 2022, 11, 98.	0.8	1