Noémie Gensous

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6428701/publications.pdf

Version: 2024-02-01

25 papers 979 citations

17 h-index

471061

25 g-index

27 all docs

 $\begin{array}{c} 27 \\ \text{docs citations} \end{array}$

times ranked

27

1997 citing authors

#	Article	IF	Citations
1	T Follicular Helper Cells in Autoimmune Disorders. Frontiers in Immunology, 2018, 9, 1637.	2.2	164
2	One-year Mediterranean diet promotes epigenetic rejuvenation with country- and sex-specific effects: a pilot study from the NU-AGE project. GeroScience, 2020, 42, 687-701.	2.1	76
3	The Impact of Caloric Restriction on the Epigenetic Signatures of Aging. International Journal of Molecular Sciences, 2019, 20, 2022.	1.8	71
4	The epigenetic landscape of age-related diseases: the geroscience perspective. Biogerontology, 2017, 18, 549-559.	2.0	62
5	Predictive biological markers of systemic lupus erythematosus flares: a systematic literature review. Arthritis Research and Therapy, 2017, 19, 238.	1.6	61
6	Immunogenicity and impact on disease activity of influenza and pneumococcal vaccines in systemic lupus erythematosus: a systematic literature review and meta-analysis. Rheumatology, 2016, 55, 1664-1672.	0.9	58
7	Multimorbidity, age-related comorbidities and mortality. Aids, 2018, 32, 1651-1660.	1.0	57
8	Age-related DNA methylation changes are sex-specific: a comprehensive assessment. Aging, 2020, 12, 24057-24080.	1.4	55
9	Accelerated bioâ€cognitive aging in Down syndrome: State of the art and possible deceleration strategies. Aging Cell, 2019, 18, e12903.	3.0	47
10	OX40L/OX40 axis impairs follicular and natural Treg function in human SLE. JCI Insight, 2018, 3, .	2.3	46
11	Responders and non-responders to influenza vaccination: A DNA methylation approach on blood cells. Experimental Gerontology, 2018, 105, 94-100.	1.2	39
12	Genomic stability, anti-inflammatory phenotype, and up-regulation of the RNAseH2 in cells from centenarians. Cell Death and Differentiation, 2019, 26, 1845-1858.	5.0	37
13	Age-Related DNA Methylation Changes: Potential Impact on Skeletal Muscle Aging in Humans. Frontiers in Physiology, 2019, 10, 996.	1.3	35
14	Down syndrome, accelerated aging and immunosenescence. Seminars in Immunopathology, 2020, 42, 635-645.	2.8	35
15	The epigenetics of inflammaging: The contribution of age-related heterochromatin loss and locus-specific remodelling and the modulation by environmental stimuli. Seminars in Immunology, 2018, 40, 49-60.	2.7	29
16	T follicular helper cells, interleukin-21 and systemic lupus erythematosus. Rheumatology, 2017, 56, kew297.	0.9	24
17	Molecular Aging of Human Liver: An Epigenetic/Transcriptomic Signature. Journals of Gerontology - Series A Biological Sciences and Medical Sciences, 2019, 74, 1-8.	1.7	23
18	Aging and Caloric Restriction Modulate the DNA Methylation Profile of the Ribosomal RNA Locus in Human and Rat Liver. Nutrients, 2020, 12, 277.	1.7	12

#	Article	IF	Citations
19	Down Syndrome, Ageing and Epigenetics. Sub-Cellular Biochemistry, 2019, 91, 161-193.	1.0	10
20	Update on the cellular pathogenesis of lupus. Current Opinion in Rheumatology, 2021, 33, 190-196.	2.0	10
21	DNA Methylation Analysis of Ribosomal DNA in Adults With Down Syndrome. Frontiers in Genetics, 2022, 13, 792165.	1.1	7
22	The impact of clopidogrel on plasma-soluble CD40 ligand levels in systemic lupus erythematosus patients: the CLOPUS phase I/II pilot study. Joint Bone Spine, 2021, 88, 105097.	0.8	3
23	Galectin 9, CXCL-10 and tumor necrosis factor receptor type II as biomarkers of disease activity in systemic lupus erythematosus. Joint Bone Spine, 2022, 89, 105311.	0.8	3
24	Cross-reactive anti-CENP-A autoantibodies induce analytic interference in anti-TIF1 \hat{I}^3 detection using line-dot immunoassay. Rheumatology, 2021, 60, 4942-4944.	0.9	2
25	GenÃ"se des anticorps. Revue Du Rhumatisme (Edition Francaise), 2016, 83, A27-A32.	0.0	0