

Liudmilla Rubbi

List of Publications by Year in Descending Order

Source: <https://exaly.com/author-pdf/6814980/liudmilla-rubbi-publications-by-year.pdf>

Version: 2024-04-19

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.

15
papers

517
citations

11
h-index

16
g-index

16
ext. papers

648
ext. citations

9.3
avg, IF

2.98
L-index

#	Paper	IF	Citations
15	Human DNA methylation signatures differentiate persistent from resolving MRSA bacteremia. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021 , 118,	11.5	3
14	Epigenetic Suppression of Transgenic T-cell Receptor Expression via Gamma-Retroviral Vector Methylation in Adoptive Cell Transfer Therapy. <i>Cancer Discovery</i> , 2020 , 10, 1645-1653	24.4	7
13	Transposon-associated epigenetic silencing during <i>Pleurotus ostreatus</i> life cycle. <i>DNA Research</i> , 2018 , 25, 451-464	4.5	13
12	Omega-3 fatty acids increase the unfolded protein response and improve amyloid- β phagocytosis by macrophages of patients with mild cognitive impairment. <i>FASEB Journal</i> , 2017 , 31, 4359-4369	0.9	18
11	Human Embryonic Stem Cells Do Not Change Their X Inactivation Status during Differentiation. <i>Cell Reports</i> , 2017 , 18, 54-67	10.6	72
10	Novel features of telomere biology revealed by the absence of telomeric DNA methylation. <i>Genome Research</i> , 2016 , 26, 1047-56	9.7	14
9	Transcriptomic Modification in the Cerebral Cortex following Noninvasive Brain Stimulation: RNA-Sequencing Approach. <i>Neural Plasticity</i> , 2016 , 2016, 5942980	3.3	4
8	The concerted impact of domestication and transposon insertions on methylation patterns between dogs and grey wolves. <i>Molecular Ecology</i> , 2016 , 25, 1838-55	5.7	52
7	Epigenetic changes in T-cell and monocyte signatures and production of neurotoxic cytokines in ALS patients. <i>FASEB Journal</i> , 2016 , 30, 3461-3473	0.9	39
6	Epigenetic changes mediated by polycomb repressive complex 2 and E2a are associated with drug resistance in a mouse model of lymphoma. <i>Genome Medicine</i> , 2016 , 8, 54	14.4	7
5	Dynamic Changes in the Transcriptome and Methylome of <i>Chlamydomonas reinhardtii</i> throughout Its Life Cycle. <i>Plant Physiology</i> , 2015 , 169, 2730-43	6.6	34
4	A high-throughput screen of inactive X chromosome reactivation identifies the enhancement of DNA demethylation by 5-aza-2edC upon inhibition of ribonucleotide reductase. <i>Epigenetics and Chromatin</i> , 2015 , 8, 42	5.8	29
3	Epigenome-wide association of liver methylation patterns and complex metabolic traits in mice. <i>Cell Metabolism</i> , 2015 , 21, 905-17	24.6	68
2	Scl binds to primed enhancers in mesoderm to regulate hematopoietic and cardiac fate divergence. <i>EMBO Journal</i> , 2015 , 34, 759-77	13	50
1	In vivo targeting of de novo DNA methylation by histone modifications in yeast and mouse. <i>ELife</i> , 2015 , 4, e06205	8.9	107