## Liudmilla Rubbi

## List of Publications by Citations

Source: https://exaly.com/author-pdf/6814980/liudmilla-rubbi-publications-by-citations.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.

16 15 517 11 h-index g-index citations papers 16 648 2.98 9.3 avg, IF L-index ext. citations ext. papers

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