Lei Huang

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8539999/publications.pdf

Version: 2024-02-01

1040056 1199594 1,079 12 9 12 citations h-index g-index papers 12 12 12 1728 all docs docs citations times ranked citing authors

7.6

13

#	Article	IF	CITATIONS
1	Mitophagy in carcinogenesis, drug resistance and anticancer therapeutics. Cancer Cell International, 2021, 21, 350.	4.1	34
2	WWP1 targeting MUC1 for ubiquitin-mediated lysosomal degradation to suppress carcinogenesis. Signal Transduction and Targeted Therapy, 2021, 6, 297.	17.1	7
3	Noninvasive preimplantation genetic testing for aneuploidy in spent medium may be more reliable than trophectoderm biopsy. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 14105-14112.	7.1	130
4	Bayesian model for accurate MARSALA (mutated allele revealed by sequencing with aneuploidy and) Tj ETQq0 0 C) rgBT /Ov 2.5	erlock 10 Tf 5
5	Multicolor Raman Beads for Multiplexed Tumor Cell and Tissue Imaging and in Vivo Tumor Spectral Detection. Analytical Chemistry, 2019, 91, 3784-3789.	6.5	45
6	Genotyping single-sperm cells by universal MARSALA enables the acquisition of linkage information for combined pre-implantation genetic diagnosis and genome screening. Journal of Assisted Reproduction and Genetics, 2018, 35, 1071-1078.	2.5	11
7	Painting a specific chromosome with CRISPR/Cas9 for live-cell imaging. Cell Research, 2017, 27, 298-301.	12.0	53
8	Single-cell whole-genome analyses by Linear Amplification via Transposon Insertion (LIANTI). Science, 2017, 356, 189-194.	12.6	303
9	Mapping allele with resolved carrier status of Robertsonian and reciprocal translocation in human preimplantation embryos. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, E8695-E8702.	7.1	36
10	Live births after simultaneous avoidance of monogenic diseases and chromosome abnormality by next-generation sequencing with linkage analyses. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 15964-15969.	7.1	115
11	Single-Cell Whole-Genome Amplification and Sequencing: Methodology and Applications. Annual Review of Genomics and Human Genetics, 2015, 16, 79-102.	6.2	327

Is there an en route folding intermediate for cold shock proteins?. Protein Science, 2012, 21, 677-685.

12