Omid R Faridani

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/2074589/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Uterine fluid microRNAs are dysregulated in women with recurrent implantation failure. Human Reproduction, 2022, 37, 734-746.	0.9	23
2	Small noncoding RNA profiling across cellular and biofluid compartments and their implications for multiple sclerosis immunopathology. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	7.1	15
3	Trophoblast derived extracellular vesicles specificallyÂalter the transcriptome of endometrial cells and may constitute a critical component of embryo-maternal communication. Reproductive Biology and Endocrinology, 2021, 19, 115.	3.3	27
4	Single-cell RNA counting at allele and isoform resolution using Smart-seq3. Nature Biotechnology, 2020, 38, 708-714.	17.5	399
5	Genomic encoding of transcriptional burst kinetics. Nature, 2019, 565, 251-254.	27.8	403
6	Small-seq for single-cell small-RNA sequencing. Nature Protocols, 2018, 13, 2407-2424.	12.0	47
7	Single-cell sequencing of the small-RNA transcriptome. Nature Biotechnology, 2016, 34, 1264-1266.	17.5	172
8	Putting cells in their place. Nature Biotechnology, 2015, 33, 490-491.	17.5	4
9	In Vivo Effects of Mesenchymal Stromal Cells in Two Patients With Severe Acute Respiratory Distress Syndrome. Stem Cells Translational Medicine, 2015, 4, 1199-1213.	3.3	131
10	Epstein–Barr virus encoded micro <scp>RNA</scp> s target <scp>SUMO</scp> â€regulated cellular functions. FEBS Journal, 2014, 281, 4935-4950.	4.7	15
11	Full-length RNA-seq from single cells using Smart-seq2. Nature Protocols, 2014, 9, 171-181.	12.0	3,308
12	Smart-seq2 for sensitive full-length transcriptome profiling in single cells. Nature Methods, 2013, 10, 1096-1098.	19.0	2,022
13	Full-length mRNA-Seq from single-cell levels of RNA and individual circulating tumor cells. Nature Biotechnology, 2012, 30, 777-782.	17.5	1,347
14	Analysis of microRNA signatures using size-coded ligation-mediated PCR. Nucleic Acids Research, 2011, 39, e80-e80.	14.5	43
15	A deneddylase encoded by Epstein–Barr virus promotes viral DNA replication by regulating the activity of cullin-RING ligases. Nature Cell Biology, 2010, 12, 351-361.	10.3	103
16	Specific ligation to double-stranded RNA for analysis of cellular RNA::RNA interactions. Nucleic Acids Research, 2008, 36, e99-e99.	14.5	7
17	Antisense PNA Accumulates in Escherichia coli and Mediates a Long Post-antibiotic Effect. Molecular Therapy, 2007, 15, 1537-1542.	8.2	59
18	Competitive inhibition of natural antisense Sok-RNA interactions activates Hok-mediated cell killing in Escherichia coli. Nucleic Acids Research, 2006, 34, 5915-5922.	14.5	60