

Lasse Ebdrup Pedersen

List of Publications by Year in descending order

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Version: 2024-02-01

26
papers

1,858
citations

471509

17
h-index

552781

26
g-index

34
all docs

34
docs citations

34
times ranked

2305
citing authors

#	ARTICLE	IF	CITATIONS
1	Rational and evolutionary engineering of <i>Saccharomyces cerevisiae</i> for production of dicarboxylic acids from lignocellulosic biomass and exploring genetic mechanisms of the yeast tolerance to the biomass hydrolysate. , 2022, 15, 22.		8
2	A synthetic RNA-mediated evolution system in yeast. <i>Nucleic Acids Research</i> , 2021, 49, e88-e88.	14.5	17
3	A metabolic CRISPR-Cas9 screen in Chinese hamster ovary cells identifies glutamine-sensitive genes. <i>Metabolic Engineering</i> , 2021, 66, 114-122.	7.0	17
4	Synergistic stabilization of a double mutant in chymotrypsin inhibitor 2 from a library screen in <i>E. coli</i> . <i>Communications Biology</i> , 2021, 4, 980.	4.4	13
5	An optimized genome-wide, virus-free CRISPR screen for mammalian cells. <i>Cell Reports Methods</i> , 2021, 1, 100062.	2.9	14
6	A dual-reporter system for investigating and optimizing protein translation and folding in <i>E. coli</i> . <i>Nature Communications</i> , 2021, 12, 6093.	12.8	12
7	Awakening dormant glycosyltransferases in CHO cells with CRISPRa. <i>Biotechnology and Bioengineering</i> , 2020, 117, 593-598.	3.3	27
8	Comprehensive Analysis of Genomic Safe Harbors as Target Sites for Stable Expression of the Heterologous Gene in HEK293 Cells. <i>ACS Synthetic Biology</i> , 2020, 9, 1263-1269.	3.8	23
9	Genome-Wide CRISPRi-Based Identification of Targets for Decoupling Growth from Production. <i>ACS Synthetic Biology</i> , 2020, 9, 1030-1040.	3.8	29
10	Multiplex secretome engineering enhances recombinant protein production and purity. <i>Nature Communications</i> , 2020, 11, 1908.	12.8	63
11	Genome-wide systematic identification of methyltransferase recognition and modification patterns. <i>Nature Communications</i> , 2019, 10, 3311.	12.8	18
12	Reprogramming AA catabolism in CHO cells with CRISPR/Cas9 genome editing improves cell growth and reduces byproduct secretion. <i>Metabolic Engineering</i> , 2019, 56, 120-129.	7.0	22
13	CasPER, a method for directed evolution in genomic contexts using mutagenesis and CRISPR/Cas9. <i>Metabolic Engineering</i> , 2018, 48, 288-296.	7.0	60
14	Ribosome profiling-guided depletion of an mRNA increases cell growth rate and protein secretion. <i>Scientific Reports</i> , 2017, 7, 40388.	3.3	48
15	Network reconstruction of the mouse secretory pathway applied on CHO cell transcriptome data. <i>BMC Systems Biology</i> , 2017, 11, 37.	3.0	14
16	A Consensus Genome-scale Reconstruction of Chinese Hamster Ovary Cell Metabolism. <i>Cell Systems</i> , 2016, 3, 434-443.e8.	6.2	205
17	CRMAGE: CRISPR Optimized MAGE Recombineering. <i>Scientific Reports</i> , 2016, 6, 19452.	3.3	180
18	CRISPy-web: An online resource to design sgRNAs for CRISPR applications. <i>Synthetic and Systems Biotechnology</i> , 2016, 1, 118-121.	3.7	117

#	ARTICLE	IF	CITATIONS
19	Accelerated homology-directed targeted integration of transgenes in Chinese hamster ovary cells via CRISPR/Cas9 and fluorescent enrichment. <i>Biotechnology and Bioengineering</i> , 2016, 113, 2518-2523.	3.3	58
20	Site-specific integration in CHO cells mediated by CRISPR/Cas9 and homology-directed DNA repair pathway. <i>Scientific Reports</i> , 2015, 5, 8572.	3.3	168
21	Multiplex metabolic pathway engineering using CRISPR/Cas9 in <i>Saccharomyces cerevisiae</i> . <i>Metabolic Engineering</i> , 2015, 28, 213-222.	7.0	355
22	One-step generation of triple knockout CHO cell lines using CRISPR/Cas9 and fluorescent enrichment. <i>Biotechnology Journal</i> , 2015, 10, 1446-1456.	3.5	108
23	Elucidation of the CHO Super-Ome (CHO-SO) by Proteoinformatics. <i>Journal of Proteome Research</i> , 2015, 14, 4687-4703.	3.7	35
24	Accelerating genome editing in CHO cells using CRISPR Cas9 and CRISPy, a web-based target finding tool. <i>Biotechnology and Bioengineering</i> , 2014, 111, 1604-1616.	3.3	167
25	High levels of the type III inorganic phosphate transporter PiT1 (SLC20A1) can confer faster cell adhesion. <i>Experimental Cell Research</i> , 2014, 326, 57-67.	2.6	20
26	Regulation of cell proliferation and cell density by the inorganic phosphate transporter PiT1. <i>Cell Division</i> , 2012, 7, 7.	2.4	39