Nicholas R Sandoval

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3230214/publications.pdf

Version: 2024-02-01

19 papers 868 citations

567281 15 h-index 19 g-index

20 all docs

20 docs citations

times ranked

20

1202 citing authors

| # | Article | IF | Citations |
|----|---|------|-----------|
| 1 | Synthetic methylotrophy: engineering the production of biofuels and chemicals based on the biology of aerobic methanol utilization. Current Opinion in Biotechnology, 2015, 33, 165-175. | 6.6 | 150 |
| 2 | Strategy for directing combinatorial genome engineering in <i>Escherichia coli</i> . Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 10540-10545. | 7.1 | 87 |
| 3 | Recent Developments of the Synthetic Biology Toolkit for Clostridium. Frontiers in Microbiology, 2018, 9, 154. | 3.5 | 85 |
| 4 | Sort-Seq Approach to Engineering a Formaldehyde-Inducible Promoter for Dynamically Regulated <i>Escherichia coli</i> Growth on Methanol. ACS Synthetic Biology, 2017, 6, 1584-1595. | 3.8 | 70 |
| 5 | Engineering membrane and cell-wall programs for tolerance to toxic chemicals: Beyond solo genes. Current Opinion in Microbiology, 2016, 33, 56-66. | 5.1 | 66 |
| 6 | Elucidating acetate tolerance in E. coli using a genome-wide approach. Metabolic Engineering, 2011, 13, 214-224. | 7.0 | 60 |
| 7 | Expression of heterologous sigma factors enables functional screening of metagenomic and heterologous genomic libraries. Nature Communications, 2015, 6, 7045. | 12.8 | 55 |
| 8 | Characterization of physiological responses to 22 gene knockouts in Escherichia coli central carbon metabolism. Metabolic Engineering, 2016, 37, 102-113. | 7.0 | 50 |
| 9 | Enzyme I facilitates reverse flux from pyruvate to phosphoenolpyruvate in Escherichia coli. Nature Communications, 2017, 8, 14316. | 12.8 | 41 |
| 10 | Transcription factor-based biosensors and inducible systems in non-model bacteria: current progress and future directions. Current Opinion in Biotechnology, 2020, 64, 39-46. | 6.6 | 37 |
| 11 | Whole-genome sequence of an evolved Clostridium pasteurianum strain reveals SpoOA deficiency responsible for increased butanol production and superior growth. Biotechnology for Biofuels, 2015, 8, 227. | 6.2 | 35 |
| 12 | Genome-Wide Mapping of Furfural Tolerance Genes in Escherichia coli. PLoS ONE, 2014, 9, e87540. | 2.5 | 30 |
| 13 | Comparison of genomeâ€wide selection strategies to identify furfural tolerance genes in <i>Escherichia coli</i> . Biotechnology and Bioengineering, 2015, 112, 129-140. | 3.3 | 30 |
| 14 | Co-utilization of glucose and xylose by evolved Thermus thermophilus LC113 strain elucidated by 13 C metabolic flux analysis and whole genome sequencing. Metabolic Engineering, 2016, 37, 63-71. | 7.0 | 27 |
| 15 | Comparison of Cross-Linked Branched and Linear Poly(ethylene imine) Microgel Microstructures and Their Impact in Antimicrobial Behavior, Copper Chelation, and Carbon Dioxide Capture. ACS Applied Polymer Materials, 2020, 2, 826-836. | 4.4 | 16 |
| 16 | Transient ammonia stress on Chinese hamster ovary (CHO) cells yield alterations to alanine metabolism and IgG glycosylation profiles. Biotechnology Journal, 2021, 16, e2100098. | 3.5 | 15 |
| 17 | Clusters of Nanoscale Liposomes Modulate the Release of Encapsulated Species and Mimic the Compartmentalization Intrinsic in Cell Structures. ACS Applied Nano Materials, 2019, 2, 7134-7143. | 5.0 | 11 |
| 18 | Synthetic biology approaches: the next tools for improved protein production from CHO cells. Current Opinion in Chemical Engineering, 2020, 30, 26-33. | 7.8 | 2 |

| # | Article | IF | CITATIONS |
|----|--|-----|-----------|
| 19 | Elucidation of Sequence–Function Relationships for an Improved Biobutanol In Vivo Biosensor in E. coli. Frontiers in Bioengineering and Biotechnology, 2022, 10, 821152. | 4.1 | 1 |