

Carlos Piñero-Lambea

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4915508/publications.pdf>

Version: 2024-02-01

10
papers

372
citations

1163117

8
h-index

1372567

10
g-index

10
all docs

10
docs citations

10
times ranked

494
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|-------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|-----|-----------|
| 1 | Programming Controlled Adhesion of <i>E. coli</i> to Target Surfaces, Cells, and Tumors with Synthetic Adhesins. ACS Synthetic Biology, 2015, 4, 463-473. | 3.8 | 133 |
| 2 | Engineered bacteria as therapeutic agents. Current Opinion in Biotechnology, 2015, 35, 94-102. | 6.6 | 83 |
| 3 | Tuning Gene Activity by Inducible and Targeted Regulation of Gene Expression in Minimal Bacterial Cells. ACS Synthetic Biology, 2018, 7, 1538-1552. | 3.8 | 30 |
| 4 | High affinity nanobodies against human epidermal growth factor receptor selected on cells by <i>E. coli</i> . MAbs, 2016, 8, 1286-1301. | 5.2 | 28 |
| 5 | <i>Mycoplasma pneumoniae</i> Genome Editing Based on Oligo Recombineering and Cas9-Mediated Counterselection. ACS Synthetic Biology, 2020, 9, 1693-1704. | 3.8 | 26 |
| 6 | Engineering a genome-reduced bacterium to eliminate <i>Staphylococcus aureus</i> biofilms <i>in vivo</i> . Molecular Systems Biology, 2021, 17, e10145. | 7.2 | 21 |
| 7 | SynMyco transposon: engineering transposon vectors for efficient transformation of minimal genomes. DNA Research, 2019, 26, 327-339. | 3.4 | 18 |
| 8 | A RAGE Based Strategy for the Genome Engineering of the Human Respiratory Pathogen <i>Mycoplasma pneumoniae</i> . ACS Synthetic Biology, 2020, 9, 2737-2748. | 3.8 | 14 |
| 9 | Inferring Active Metabolic Pathways from Proteomics and Essentiality Data. Cell Reports, 2020, 31, 107722. | 6.4 | 10 |
| 10 | LoxTnSeq: random transposon insertions combined with cre/lox recombination and counterselection to generate large random genome reductions. Microbial Biotechnology, 2021, 14, 2403-2419. | 4.2 | 9 |