

# Daniel B Goodman

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8980890/publications.pdf>

Version: 2024-02-01

13  
papers

2,599  
citations

840776

11  
h-index

1125743

13  
g-index

18  
all docs

18  
docs citations

18  
times ranked

3134  
citing authors

| #  | ARTICLE   | IF   | CITATIONS |
|----|---|------|-----------|
| 1  | High-throughput functional variant screens via in vivo production of single-stranded DNA. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .                               | 7.1  | 53        |
| 2  | Enabling multiplexed testing of pooled donor cells through whole-genome sequencing. Genome Medicine, 2018, 10, 31.  | 8.2  | 10        |
| 3  | Optimizing complex phenotypes through model-guided multiplex genome engineering. Genome Biology, 2017, 18, 100.   | 8.8  | 23        |
| 4  | Millstone: software for multiplex microbial genome analysis and engineering. Genome Biology, 2017, 18, 101.   | 8.8  | 5         |
| 5  | DNAplotlib: Programmable Visualization of Genetic Designs and Associated Data. ACS Synthetic Biology, 2017, 6, 1115-1119.   | 3.8  | 50        |
| 6  | Emergent rules for codon choice elucidated by editing rare arginine codons in <i>Escherichia coli</i> . Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, E5588-97.         | 7.1  | 48        |
| 7  | Design, synthesis, and testing toward a 57-codon genome. Science, 2016, 353, 819-822.   | 12.6 | 251       |
| 8  | Rational optimization of <i>tolC</i> as a powerful dual selectable marker for genome engineering. Nucleic Acids Research, 2014, 42, 4779-4790.  | 14.5 | 36        |
| 9  | Composability of regulatory sequences controlling transcription and translation in <i>Escherichia coli</i> . Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 14024-14029. | 7.1  | 377       |
| 10 | Genomically Recoded Organisms Expand Biological Functions. Science, 2013, 342, 357-360.   | 12.6 | 721       |
| 11 | Causes and Effects of N-Terminal Codon Bias in Bacterial Genes. Science, 2013, 342, 475-479.  | 12.6 | 491       |
| 12 | Precise Manipulation of Chromosomes in Vivo Enables Genome-Wide Codon Replacement. Science, 2011, 333, 348-353.   | 12.6 | 512       |
| 13 | Product Length, Dye Choice, and Detection Chemistry in the Bead-Emulsion Amplification of Millions of Single DNA Molecules in Parallel. Analytical Chemistry, 2009, 81, 5770-5776.                                    | 6.5  | 15        |