## David Botstein

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/9719754/publications.pdf

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759233 1125743 1,833 13 12 13 citations h-index g-index papers 14 14 14 2873 docs citations times ranked citing authors all docs

| #  | Article   | lF   | CITATIONS |
|----|---|------|-----------|
| 1  | Coordination of Growth Rate, Cell Cycle, Stress Response, and Metabolic Activity in Yeast. Molecular Biology of the Cell, 2008, 19, 352-367.  | 2.1  | 524       |
| 2  | Yeast: An Experimental Organism for 21st Century Biology. Genetics, 2011, 189, 695-704.   | 2.9  | 450       |
| 3  | A molecular barcoded yeast ORF library enables mode-of-action analysis of bioactive compounds.<br>Nature Biotechnology, 2009, 27, 369-377.  | 17.5 | 254       |
| 4  | Synthetic gene expression perturbation systems with rapid, tunable, single-gene specificity in yeast. Nucleic Acids Research, 2013, 41, e57-e57.  | 14.5 | 141       |
| 5  | Fast-acting and nearly gratuitous induction of gene expression and protein depletion in <i>Saccharomyces cerevisiae</i> Molecular Biology of the Cell, 2011, 22, 4447-4459.   | 2.1  | 120       |
| 6  | Synthetic biology tools for programming gene expression without nutritional perturbations in Saccharomyces cerevisiae. Nucleic Acids Research, 2014, 42, e48-e48.   | 14.5 | 87        |
| 7  | Characterizing the in vivo role of trehalose in <i>Saccharomyces cerevisiae</i> using the <i>AGT1</i> transporter. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 6116-6121. | 7.1  | 77        |
| 8  | Perturbation-based analysis and modeling of combinatorial regulation in the yeast sulfur assimilation pathway. Molecular Biology of the Cell, 2012, 23, 2993-3007.  | 2.1  | 45        |
| 9  | Novel insights from a multiomics dissection of the Hayflick limit. ELife, 2022, 11, .   | 6.0  | 38        |
| 10 | A genomeâ€scale yeast library with inducible expression of individual genes. Molecular Systems Biology, 2021, 17, e10207.   | 7.2  | 37        |
| 11 | The future of humans as model organisms. Science, 2018, 361, 552-553.   | 12.6 | 31        |
| 12 | Simultaneous Profiling of DNA Accessibility and Gene Expression Dynamics with ATAC-Seq and RNA-Seq. Methods in Molecular Biology, 2018, 1819, 317-333.  | 0.9  | 18        |
| 13 | Rapid Synthesis and Screening of Chemically Activated Transcription Factors with GFP-based Reporters. Journal of Visualized Experiments, 2013, , e51153.  | 0.3  | 11        |