

Mark P Styczynski

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3103833/publications.pdf>

Version: 2024-02-01

51
papers

1,416
citations

430754

18
h-index

345118

36
g-index

59
all docs

59
docs citations

59
times ranked

2270
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|-----|-----------|
| 1 | Systematic Identification of Conserved Metabolites in GC/MS Data for Metabolomics and Biomarker Discovery. <i>Analytical Chemistry</i> , 2007, 79, 966-973. | 3.2 | 223 |
| 2 | The Cancer Microbiome: Distinguishing Direct and Indirect Effects Requires a Systemic View. <i>Trends in Cancer</i> , 2020, 6, 192-204. | 3.8 | 162 |
| 3 | Applications of metabolomics in cancer research. <i>Journal of Carcinogenesis</i> , 2013, 12, 9. | 2.5 | 83 |
| 4 | BLOSUM62 miscalculations improve search performance. <i>Nature Biotechnology</i> , 2008, 26, 274-275. | 9.4 | 79 |
| 5 | Point-of-care biomarker quantification enabled by sample-specific calibration. <i>Science Advances</i> , 2019, 5, eaax4473. | 4.7 | 73 |
| 6 | A Multi-Modality CMOS Sensor Array for Cell-Based Assay and Drug Screening. <i>IEEE Transactions on Biomedical Circuits and Systems</i> , 2015, 9, 801-814. | 2.7 | 65 |
| 7 | Development of a Pigment-Based Whole-Cell Zinc Biosensor for Human Serum. <i>ACS Synthetic Biology</i> , 2018, 7, 267-275. | 1.9 | 59 |
| 8 | Precision metabolic engineering: The design of responsive, selective, and controllable metabolic systems. <i>Metabolic Engineering</i> , 2015, 31, 123-131. | 3.6 | 56 |
| 9 | Overview of computational methods for the inference of gene regulatory networks. <i>Computers and Chemical Engineering</i> , 2005, 29, 519-534. | 2.0 | 48 |
| 10 | NS-kNN: a modified k-nearest neighbors approach for imputing metabolomics data. <i>Metabolomics</i> , 2018, 14, 153. | 1.4 | 44 |
| 11 | Metabolic Profiling of <i>Escherichia coli</i> -Based Cell-Free Expression Systems for Process Optimization. <i>Industrial & Engineering Chemistry Research</i> , 2019, 58, 22472-22482. | 1.8 | 30 |
| 12 | OVCAR-3 Spheroid-Derived Cells Display Distinct Metabolic Profiles. <i>PLoS ONE</i> , 2015, 10, e0118262. | 1.1 | 29 |
| 13 | Precise metabolic engineering of carotenoid biosynthesis in <i>Escherichia coli</i> towards a low-cost biosensor. <i>Metabolic Engineering</i> , 2015, 31, 171-180. | 3.6 | 28 |
| 14 | Dynamic and tunable metabolite control for robust minimal-equipment assessment of serum zinc. <i>Nature Communications</i> , 2019, 10, 5514. | 5.8 | 26 |
| 15 | Metabolomics identifies the intersection of phosphoethanolamine with menaquinone-triggered apoptosis in an in vitro model of leukemia. <i>Molecular BioSystems</i> , 2015, 11, 2406-2416. | 2.9 | 25 |
| 16 | Distinct metabolic responses of an ovarian cancer stem cell line. <i>BMC Systems Biology</i> , 2014, 8, 134. | 3.0 | 23 |
| 17 | Precise control of lycopene production to enable a fast-responding, minimal-equipment biosensor. <i>Metabolic Engineering</i> , 2017, 43, 46-53. | 3.6 | 23 |
| 18 | Active Analyte Import Improves the Dynamic Range and Sensitivity of a Vitamin B ₁₂ Biosensor. <i>ACS Synthetic Biology</i> , 2020, 9, 402-411. | 1.9 | 22 |

| # | ARTICLE | IF | CITATIONS |
|----|--|-----|-----------|
| 19 | Systematic Applications of Metabolomics in Metabolic Engineering. <i>Metabolites</i> , 2012, 2, 1090-1122. | 1.3 | 20 |
| 20 | Integrative analysis associates monocytes with insufficient erythropoiesis during acute <i>Plasmodium cynomolgi</i> malaria in rhesus macaques. <i>Malaria Journal</i> , 2017, 16, 384. | 0.8 | 20 |
| 21 | Systems Biology-Based Investigation of Host-Plasmodium Interactions. <i>Trends in Parasitology</i> , 2018, 34, 617-632. | 1.5 | 19 |
| 22 | Metabolomics Analysis of the Toxic Effects of the Production of Lycopene and Its Precursors. <i>Frontiers in Microbiology</i> , 2018, 9, 760. | 1.5 | 19 |
| 23 | Metabolic modeling helps interpret transcriptomic changes during malaria. <i>Biochimica Et Biophysica Acta - Molecular Basis of Disease</i> , 2018, 1864, 2329-2340. | 1.8 | 18 |
| 24 | Protocell arrays for simultaneous detection of diverse analytes. <i>Nature Communications</i> , 2021, 12, 5724. | 5.8 | 18 |
| 25 | Humoral immunity prevents clinical malaria during <i>Plasmodium</i> relapses without eliminating gametocytes. <i>PLoS Pathogens</i> , 2019, 15, e1007974. | 2.1 | 17 |
| 26 | ElectroPen: An ultra-low-cost, electricity-free, portable electroporator. <i>PLoS Biology</i> , 2020, 18, e3000589. | 2.6 | 16 |
| 27 | Comparative transcriptomics and metabolomics in a rhesus macaque drug administration study. <i>Frontiers in Cell and Developmental Biology</i> , 2014, 2, 54. | 1.8 | 15 |
| 28 | Effective Use of Linear DNA in Cell-Free Expression Systems. <i>Frontiers in Bioengineering and Biotechnology</i> , 2021, 9, 715328. | 2.0 | 15 |
| 29 | Small molecule signaling, regulation, and potential applications in cellular therapeutics. <i>Wiley Interdisciplinary Reviews: Systems Biology and Medicine</i> , 2018, 10, e1405. | 6.6 | 14 |
| 30 | Differential metabolite levels in response to spawning-induced inappetence in Atlantic salmon <i>Salmo salar</i> . <i>Comparative Biochemistry and Physiology Part D: Genomics and Proteomics</i> , 2015, 13, 52-59. | 0.4 | 13 |
| 31 | Metabolic Dynamics in <i>Escherichia coli</i> -Based Cell-Free Systems. <i>ACS Synthetic Biology</i> , 2021, 10, 2252-2265. | 1.9 | 11 |
| 32 | An extension and novel solution to the (l,d)-motif challenge problem. <i>Genome Informatics</i> , 2004, 15, 63-71. | 0.4 | 11 |
| 33 | Reliable computation of equilibrium states and bifurcations in food chain models. <i>Computers and Chemical Engineering</i> , 2004, 28, 1981-1996. | 2.0 | 9 |
| 34 | Low-cost, point-of-care biomarker quantification. <i>Current Opinion in Biotechnology</i> , 2022, 76, 102738. | 3.3 | 9 |
| 35 | From genome-scale data to models of infectious disease: A Bayesian network-based strategy to drive model development. <i>Mathematical Biosciences</i> , 2015, 270, 156-168. | 0.9 | 8 |
| 36 | LK-DFBA: a linear programming-based modeling strategy for capturing dynamics and metabolite-dependent regulation in metabolism. <i>BMC Bioinformatics</i> , 2020, 21, 93. | 1.2 | 7 |

| # | ARTICLE | IF | CITATIONS |
|----|---|-----|-----------|
| 37 | A tree-like Bayesian structure learning algorithm for small-sample datasets from complex biological model systems. <i>BMC Systems Biology</i> , 2015, 9, 49. | 3.0 | 6 |
| 38 | Improved metabolite profile smoothing for flux estimation. <i>Molecular BioSystems</i> , 2015, 11, 2394-2405. | 2.9 | 6 |
| 39 | Point-of-Care Analyte Quantification and Digital Readout <i>via</i> Lysate-Based Cell-Free Biosensors Interfaced with Personal Glucose Monitors. <i>ACS Synthetic Biology</i> , 2021, 10, 2862-2869. | 1.9 | 6 |
| 40 | Dramatic transcriptomic differences in <i>Macaca mulatta</i> and <i>Macaca fascicularis</i> with <i>Plasmodium knowlesi</i> infections. <i>Scientific Reports</i> , 2021, 11, 19519. | 1.6 | 5 |
| 41 | Manipulation of metabolism in complex eukaryotic systems to control cellular state. <i>Current Opinion in Chemical Engineering</i> , 2015, 10, 63-69. | 3.8 | 4 |
| 42 | Systems biology-based analysis of cell-free systems. <i>Current Opinion in Biotechnology</i> , 2022, 75, 102703. | 3.3 | 4 |
| 43 | The Metabolomics Society's Current State of the Membership and Future Directions. <i>Metabolites</i> , 2019, 9, 89. | 1.3 | 2 |
| 44 | SCOUR: a stepwise machine learning framework for predicting metabolite-dependent regulatory interactions. <i>BMC Bioinformatics</i> , 2021, 22, 365. | 1.2 | 2 |
| 45 | Live demonstration: A 1024-pixel CMOS multi-modality sensing array for cell-based assays. , 2016, , . | | 1 |
| 46 | Gas Chromatography-Mass Spectrometry Microbial Metabolomics for Applications in Strain Optimization. <i>Methods in Molecular Biology</i> , 2019, 1927, 179-189. | 0.4 | 1 |
| 47 | Untargeted plasma metabolomic analysis of wild bottlenose dolphins (<i>Tursiops truncatus</i>) indicate protein degradation when in poorer health. <i>Comparative Biochemistry and Physiology Part D: Genomics and Proteomics</i> , 2022, 42, 100991. | 0.4 | 1 |
| 48 | Nucleic Acid Partitioning in PEG-Ficoll Protocells. <i>Journal of Chemical & Engineering Data</i> , 2022, 67, 1964-1971. | 1.0 | 1 |
| 49 | Editorial overview: Systems biology-the intersection of experiments and computation, underpinning biotechnology. <i>Current Opinion in Biotechnology</i> , 2016, 39, iv-vi. | 3.3 | 0 |
| 50 | Diverse classes of constraints enable broader applicability of a linear programming-based dynamic metabolic modeling framework. <i>Scientific Reports</i> , 2022, 12, 762. | 1.6 | 0 |
| 51 | Metabolomics Analysis of Cell-Free Systems Using. <i>Methods in Molecular Biology</i> , 2022, 2433, 217-226. | 0.4 | 0 |