## Michael Eisenstein

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/10819097/publications.pdf

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42 papers

2,099 citations

236925 25 h-index 243625 44 g-index

55 all docs 55 docs citations

55 times ranked 3207 citing authors

#	Article	IF	CITATIONS
1	Real-Time, Aptamer-Based Tracking of Circulating Therapeutic Agents in Living Animals. Science Translational Medicine, 2013, 5, 213ra165.	12.4	291
2	Oxford Nanopore announcement sets sequencing sector abuzz. Nature Biotechnology, 2012, 30, 295-296.	17.5	156
3	Integrated Electrochemical Microsystems for Genetic Detection of Pathogens at the Point of Care. Accounts of Chemical Research, 2015, 48, 911-920.	15.6	135
4	Quantitative selection and parallel characterization of aptamers. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 18460-18465.	7.1	115
5	Synthetic Aptamer-Polymer Hybrid Constructs for Programmed Drug Delivery into Specific Target Cells. Journal of the American Chemical Society, 2014, 136, 15010-15015.	13.7	110
6	Particle Display: A Quantitative Screening Method for Generating Highâ€Affinity Aptamers. Angewandte Chemie - International Edition, 2014, 53, 4796-4801.	13.8	96
7	A Cellular Repressor of E1A-Stimulated Genes That Inhibits Activation by E2F. Molecular and Cellular Biology, 1998, 18, 5032-5041.	2.3	87
8	The secreted glycoprotein CREG enhances differentiation of NTERA-2 human embryonal carcinoma cells. Oncogene, 2000, 19, 2120-2128.	5.9	76
9	Multiparameter Particle Display (MPPD): A Quantitative Screening Method for the Discovery of Highly Specific Aptamers. Angewandte Chemie - International Edition, 2017, 56, 744-747.	13.8	71
10	Engineering Aptamer Switches for Multifunctional Stimulusâ€Responsive Nanosystems. Advanced Materials, 2020, 32, e2003704.	21.0	68
11	Dual-reporter SERS-based biomolecular assay with reduced false-positive signals. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, 9056-9061.	7.1	67
12	Independent control of the thermodynamic and kinetic properties of aptamer switches. Nature Communications, 2019, 10, 5079.	12.8	62
13	High-Fidelity Nanopore Sequencing of Ultra-Short DNA Targets. Analytical Chemistry, 2019, 91, 6783-6789.	6.5	50
14	Phenotypic effects of an induced mutation of the ObRa isoform of the leptin receptor. Molecular Metabolism, 2013, 2, 364-375.	6.5	49
15	Startups use short-read data to expand long-read sequencing market. Nature Biotechnology, 2015, 33, 433-435.	17.5	48
16	Rational design of aptamer switches with programmable pH response. Nature Communications, 2020, 11, 2946.	12.8	45
17	A fluorescence sandwich immunoassay for the real-time continuous detection of glucose and insulin in live animals. Nature Biomedical Engineering, 2021, 5, 53-63.	22.5	44
18	Direct Selection of Fluorescence-Enhancing RNA Aptamers. Journal of the American Chemical Society, 2018, 140, 3583-3591.	13.7	42

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19	Thousand-Fold Volumetric Concentration of Live Cells with a Recirculating Acoustofluidic Device. Analytical Chemistry, 2015, 87, 8497-8502.	6.5	39
20	Array-based Discovery of Aptamer Pairs. Analytical Chemistry, 2015, 87, 821-828.	<b>6.</b> 5	39
21	Directed Evolution of Aptamer Discovery Technologies. Accounts of Chemical Research, 2022, 55, 685-695.	15.6	35
22	Accelerated Electron Transfer in Nanostructured Electrodes Improves the Sensitivity of Electrochemical Biosensors. Advanced Science, 2021, 8, e2102495.	11.2	32
23	Accurate Zygoteâ€Specific Discrimination of Singleâ€Nucleotide Polymorphisms Using Microfluidic Electrochemical DNA Melting Curves. Angewandte Chemie - International Edition, 2014, 53, 3163-3167.	13.8	29
24	Shape-based separation of synthetic microparticles. Nature Materials, 2019, 18, 82-89.	27.5	29
25	Discovery of indole-modified aptamers for highly specific recognition of protein glycoforms. Nature Communications, 2021, 12, 7106.	12.8	28
26	<i>In Vitro</i> Selection of Shape-Changing DNA Nanostructures Capable of Binding-Induced Cargo Release. ACS Nano, 2013, 7, 9675-9683.	14.6	26
27	Real-time monitoring of drug pharmacokinetics within tumor tissue in live animals. Science Advances, 2022, 8, eabk2901.	10.3	26
28	The field that came in from the cold. Nature Methods, 2016, 13, 19-22.	19.0	23
29	The battle for sequencing supremacy. Nature Biotechnology, 2012, 30, 1023-1026.	17.5	20
30	A system for multiplexed selection of aptamers with exquisite specificity without counterselection. Proceedings of the National Academy of Sciences of the United States of America, 2022, 119, e2119945119.	7.1	20
31	Direct Selection Strategy for Isolating Aptamers with pH-Sensitive Binding Activity. ACS Sensors, 2018, 3, 2574-2580.	7.8	17
32	High-Throughput Discovery of Aptamers for Sandwich Assays. Analytical Chemistry, 2016, 88, 10842-10847.	6.5	14
33	Measuring Aptamer Folding Energy Using a Molecular Clamp. Journal of the American Chemical Society, 2020, 142, 11743-11749.	13.7	9
34	Up for grabs. Nature Biotechnology, 2010, 28, 544-546.	17.5	7
35	Personalized, sequencing-based immune profiling spurs startups. Nature Biotechnology, 2013, 31, 184-185.	17.5	7
36	Strategy for Generating Sequence-Defined Aptamer Reagent Sets for Detecting Protein Contaminants in Biotherapeutics. Analytical Chemistry, 2018, 90, 3262-3269.	6.5	7

#	Article	IF	CITATIONS
37	Companies 'going long' generate sequencing buzz at Marco Island. Nature Biotechnology, 2013, 31, 265-266.	17.5	6
38	Multiparameter Particle Display (MPPD): A Quantitative Screening Method for the Discovery of Highly Specific Aptamers. Angewandte Chemie, 2017, 129, 762-765.	2.0	6
39	Illumina swallows PacBio in long shot for market domination. Nature Biotechnology, 2019, 37, 3-4.	17.5	4
40	Comparing assays via the resolution of molecular concentration. Nature Biomedical Engineering, 2022, 6, 227-231.	22.5	4
41	Immune profiling players shift gear to guide cancer drug development. Nature Biotechnology, 2016, 34, 215-216.	17.5	1
42	Frontispiece: Particle Display: A Quantitative Screening Method for Generating High-Affinity Aptamers. Angewandte Chemie - International Edition, 2014, 53, n/a-n/a.	13.8	0