

# Eldon Emberly

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/60251/publications.pdf>

Version: 2024-02-01

33  
papers

1,798  
citations

566801

15  
h-index

454577

30  
g-index

33  
all docs

33  
docs citations

33  
times ranked

3694  
citing authors

#	ARTICLE	IF	CITATIONS
1	Optimizing Efficiency and Motility of a Polyvalent Molecular Motor. <i>Micromachines</i> , 2022, 13, 914.	1.4	0
2	Substrate stiffness tunes the dynamics of polyvalent rolling motors. <i>Soft Matter</i> , 2021, 17, 1468-1479.	1.2	20
3	Functional mapping of androgen receptor enhancer activity. <i>Genome Biology</i> , 2021, 22, 149.	3.8	18
4	Abrupt, Asynchronous Changes in Action Representations by Anterior Cingulate Cortex Neurons during Trial and Error Learning. <i>Cerebral Cortex</i> , 2020, 30, 4336-4345.	1.6	4
5	DNA segregation under Par protein control. <i>PLoS ONE</i> , 2019, 14, e0218520.	1.1	9
6	Children's biobehavioral reactivity to challenge predicts DNA methylation in adolescence and emerging adulthood. <i>Developmental Science</i> , 2019, 22, e12739.	1.3	6
7	Dense neural networks for predicting chromatin conformation. <i>BMC Bioinformatics</i> , 2018, 19, 372.	1.2	13
8	A maximum-entropy model for predicting chromatin contacts. <i>PLoS Computational Biology</i> , 2018, 14, e1005956.	1.5	9
9	Confinement-dependent localization of diffusing aggregates in cellular geometries. <i>Physical Review E</i> , 2015, 91, 012705.	0.8	0
10	Bacterial motion in narrow capillaries. <i>FEMS Microbiology Ecology</i> , 2015, 91, 1-7.	1.3	7
11	Concordant and discordant DNA methylation signatures of aging in human blood and brain. <i>Epigenetics and Chromatin</i> , 2015, 8, 19.	1.8	132
12	Probing long-range interactions by extracting free energies from genome-wide chromosome conformation capture data. <i>BMC Bioinformatics</i> , 2015, 16, 171.	1.2	3
13	Operational Principles for the Dynamics of the In Vitro ParA-ParB System. <i>PLoS Computational Biology</i> , 2015, 11, e1004651.	1.5	18
14	Localization of aggregating proteins in bacteria depends on the rate of addition. <i>Frontiers in Microbiology</i> , 2014, 5, 418.	1.5	8
15	Chromatin Immunoprecipitation Indirect Peaks Highlight Long-Range Interactions of Insulator Proteins and Pol II Pausing. <i>Molecular Cell</i> , 2014, 53, 672-681.	4.5	102
16	Insulators recruit histone methyltransferase <i>Mes4</i> to regulate chromatin of flanking genes. <i>EMBO Journal</i> , 2014, 33, 1599-1613.	3.5	34
17	Additional annotation enhances potential for biologically-relevant analysis of the Illumina Infinium HumanMethylation450 BeadChip array. <i>Epigenetics and Chromatin</i> , 2013, 6, 4.	1.8	412
18	Reply to Suderman et al.: Importance of accounting for blood cell composition in epigenetic studies. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, E1247.	3.3	7

#	ARTICLE	IF	CITATIONS
19	Non-Equilibrium Polar Localization of Proteins in Bacterial Cells. PLoS ONE, 2013, 8, e64075.	1.1	7
20	A Model for Cell Population Size Control Using Asymmetric Division. PLoS ONE, 2013, 8, e74324.	1.1	0
21	CHROMATRA: a Galaxy tool for visualizing genome-wide chromatin signatures. Bioinformatics, 2012, 28, 717-718.	1.8	14
22	Factors underlying variable DNA methylation in a human community cohort. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 17253-17260.	3.3	414
23	Vital Dye Reaction and Granule Localization in Periplasm of Escherichia coli. PLoS ONE, 2012, 7, e38427.	1.1	10
24	Splitting the task: Ubp8 and Ubp10 deubiquitinate different cellular pools of H2BK123. Genes and Development, 2011, 25, 2242-2247.	2.7	96
25	Chromosome Driven Spatial Patterning of Proteins in Bacteria. PLoS Computational Biology, 2010, 6, e1000986.	1.5	32
26	Genome-Wide Mapping of Boundary Element-Associated Factor (BEAF) Binding Sites in <i>Drosophila melanogaster</i> Links BEAF to Transcription. Molecular and Cellular Biology, 2009, 29, 3556-3568.	1.1	95
27	Optimizing the readout of morphogen gradients. Physical Review E, 2008, 77, 041903.	0.8	13
28	BEAF Regulates Cell-Cycle Genes through the Controlled Deposition of H3K9 Methylation Marks into Its Conserved Dual-Core Binding Sites. PLoS Biology, 2008, 6, e327.	2.6	60
29	Hourglass Model for a Protein-Based Circadian Oscillator. Physical Review Letters, 2006, 96, 038303.	2.9	59
30	Conservation of regulatory elements between two species of <i>Drosophila</i> . BMC Bioinformatics, 2003, 4, 57.	1.2	84
31	Principles for the design and operation of a molecular wire transistor. Journal of Applied Physics, 2000, 88, 5280-5282.	1.1	18
32	Electrical conductance of molecular wires. Nanotechnology, 1999, 10, 285-289.	1.3	37
33	State Orthogonalization by Building a Hilbert Space: A New Approach to Electronic Quantum Transport in Molecular Wires. Physical Review Letters, 1998, 81, 5205-5208.	2.9	57