

# Daniel D Seaton

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4895839/publications.pdf>

Version: 2024-02-01

17  
papers

1,090  
citations

567144

15  
h-index

887953

17  
g-index

26  
all docs

26  
docs citations

26  
times ranked

1950  
citing authors

#	ARTICLE	IF	CITATIONS
1	Identification of rare and common regulatory variants in pluripotent cells using population-scale transcriptomics. <i>Nature Genetics</i> , 2021, 53, 313-321.	9.4	42
2	Population-scale single-cell RNA-seq profiling across dopaminergic neuron differentiation. <i>Nature Genetics</i> , 2021, 53, 304-312.	9.4	146
3	Discovery and quality analysis of a comprehensive set of structural variants and short tandem repeats. <i>Nature Communications</i> , 2020, 11, 2928.	5.8	22
4	Single-cell RNA-sequencing of differentiating iPS cells reveals dynamic genetic effects on gene expression. <i>Nature Communications</i> , 2020, 11, 810.	5.8	235
5	Population-scale proteome variation in human induced pluripotent stem cells. <i>ELife</i> , 2020, 9, .	2.8	40
6	An explanatory model of temperature influence on flowering through whole-plant accumulation of FLOWERING LOCUS T in <i>Arabidopsis thaliana</i> . <i>In Silico Plants</i> , 2019, 1, .	0.8	20
7	Photoperiodic control of the <i>Arabidopsis</i> proteome reveals a translational coincidence mechanism. <i>Molecular Systems Biology</i> , 2018, 14, e7962.	3.2	74
8	Dawn and photoperiod sensing by phytochrome A. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 10523-10528.	3.3	34
9	Multi-scale modelling to synergise Plant Systems Biology and Crop Science. <i>Field Crops Research</i> , 2017, 202, 77-83.	2.3	21
10	ODE-Based Modeling of Complex Regulatory Circuits. <i>Methods in Molecular Biology</i> , 2017, 1629, 317-330.	0.4	0
11	Model-Based Analysis of Cell Cycle Responses to Dynamically Changing Environments. <i>PLoS Computational Biology</i> , 2016, 12, e1004604.	1.5	12
12	Photoperiod-dependent changes in the phase of core clock transcripts and global transcriptional outputs at dawn and dusk in <i>Arabidopsis</i> . <i>Plant, Cell and Environment</i> , 2016, 39, 1955-1981.	2.8	60
13	Photoreceptor effects on plant biomass, resource allocation, and metabolic state. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 7667-7672.	3.3	115
14	Defining the robust behaviour of the plant clock gene circuit with absolute RNA timeseries and open infrastructure. <i>Open Biology</i> , 2015, 5, 150042.	1.5	42
15	Linked circadian outputs control elongation growth and flowering in response to photoperiod and temperature. <i>Molecular Systems Biology</i> , 2015, 11, 776.	3.2	87
16	Regulatory principles and experimental approaches to the circadian control of starch turnover. <i>Journal of the Royal Society Interface</i> , 2014, 11, 20130979.	1.5	29
17	Mathematical Models Light Up Plant Signaling. <i>Plant Cell</i> , 2014, 26, 5-20.	3.1	41