## Mathew G Lewsey

## List of Publications by Year in Descending Order

Source: https://exaly.com/author-pdf/3655669/mathew-g-lewsey-publications-by-year.pdf

Version: 2024-04-17

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

45
papers

4,096
citations

h-index

58
g-index

58
ext. papers

9.9
avg, IF

L-index

#	Paper	IF	Citations
45	scCloudMine: A cloud-based app for visualization, comparison, and exploration of single cell transcriptomic data. <i>Plant Communications</i> , <b>2022</b> , 100302	9	
44	Insights into opium poppy (Papaver spp.) genetic diversity from genotyping-by-sequencing analysis <i>Scientific Reports</i> , <b>2022</b> , 12, 111	4.9	0
43	Applications of hyperspectral imaging in plant phenotyping Trends in Plant Science, 2022,	13.1	5
42	Applications of cell- and tissue-specific <b>V</b> omics to improve plant productivity <i>Emerging Topics in Life Sciences</i> , <b>2022</b> ,	3.5	1
41	How Vomics technologies can drive plant engineering, ecosystem surveillance, human and animal health <i>Emerging Topics in Life Sciences</i> , <b>2022</b> , 6, 137-139	3.5	
40	RNA-seq analysis of laser microdissected Arabidopsis thaliana leaf epidermis, mesophyll and vasculature defines tissue-specific transcriptional responses to multiple stress treatments. <i>Plant Journal</i> , <b>2021</b> , 107, 938-955	6.9	6
39	Noninvasive imaging technologies in plant phenotyping. <i>Trends in Plant Science</i> , <b>2021</b> ,	13.1	2
38	Recent advances in Cannabis sativa genomics research. New Phytologist, 2021, 230, 73-89	9.8	23
37	Small RNAs shoot for the root. <i>Nature Plants</i> , <b>2021</b> , 7, 2-3	11.5	1
36	Vision, challenges and opportunities for a Plant Cell Atlas. <i>ELife</i> , <b>2021</b> , 10,	8.9	8
35	Analysis of Spatio-Temporal Transcriptome Profiles of Soybean () Tissues during Early Seed Development. <i>International Journal of Molecular Sciences</i> , <b>2020</b> , 21,	6.3	3
34	Integrated multi-omics framework of the plant response to jasmonic acid. <i>Nature Plants</i> , <b>2020</b> , 6, 290-3	<b>02</b> 1.5	59
33	Laser-Capture Microdissection RNA-Sequencing for Spatial and Temporal Tissue-Specific Gene Expression Analysis in Plants. <i>Journal of Visualized Experiments</i> , <b>2020</b> ,	1.6	1
32	Plant Pathology 101: how to get away with infection. New Phytologist, 2020, 225, 601-603	9.8	0
31	Temporal tissue-specific regulation of transcriptomes during barley (Hordeum vulgare) seed germination. <i>Plant Journal</i> , <b>2020</b> , 101, 700-715	6.9	8
30	The JA-pathway MYC transcription factors regulate photomorphogenic responses by targeting HY5 gene expression. <i>Plant Journal</i> , <b>2020</b> , 102, 138-152	6.9	24
29	Developmental normalization of phenomics data generated by high throughput plant phenotyping systems. <i>Plant Methods</i> , <b>2020</b> , 16, 111	5.8	O

## (2011-2019)

28	ANAC017 Coordinates Organellar Functions and Stress Responses by Reprogramming Retrograde Signaling. <i>Plant Physiology</i> , <b>2019</b> , 180, 634-653	6.6	36
27	A MYC2/MYC3/MYC4-dependent transcription factor network regulates water spray-responsive gene expression and jasmonate levels. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2019</b> , 116, 23345-23356	11.5	38
26	Epigenetic silencing of a multifunctional plant stress regulator. ELife, 2019, 8,	8.9	16
25	Biochemistry, Genetics, and Genomics of Opium Poppy (Papaver somniferum) for Crop Improvement <b>2019</b> , 1177-1219		2
24	Regulation of genome-wide DNA methylation by mobile small RNAs. New Phytologist, 2018, 217, 540-5	<b>46</b> .8	38
23	AgriSeqDB: an online RNA-Seq database for functional studies of agriculturally relevant plant species. <i>BMC Plant Biology</i> , <b>2018</b> , 18, 200	5.3	7
22	JAZ2 controls stomata dynamics during bacterial invasion. New Phytologist, 2017, 213, 1378-1392	9.8	80
21	Extensive transcriptomic and epigenomic remodelling occurs during Arabidopsis thaliana germination. <i>Genome Biology</i> , <b>2017</b> , 18, 172	18.3	87
20	Salicylic acid treatment and expression of an RNA-dependent RNA polymerase 1 transgene inhibit lethal symptoms and meristem invasion during tobacco mosaic virus infection in Nicotiana benthamiana. <i>BMC Plant Biology</i> , <b>2016</b> , 16, 15	5.3	42
19	Mobile small RNAs regulate genome-wide DNA methylation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2016</b> , 113, E801-10	11.5	153
18	Cistrome and Epicistrome Features Shape the Regulatory DNA Landscape. <i>Cell</i> , <b>2016</b> , 165, 1280-1292	56.2	528
17	Mobile small RNAs and their role in regulating cytosine methylation of DNA. RNA Biology, 2016, 13, 106	504.1806	7 5
16	Domains of the cucumber mosaic virus 2b silencing suppressor protein affecting inhibition of salicylic acid-induced resistance and priming of salicylic acid accumulation during infection. <i>Journal of General Virology</i> , <b>2014</b> , 95, 1408-1413	4.9	30
15	Interference with jasmonic acid-regulated gene expression is a general property of viral suppressors of RNA silencing but only partly explains virus-induced changes in plant-aphid interactions. <i>Journal of General Virology</i> , <b>2014</b> , 95, 733-739	4.9	46
14	Determination and inference of eukaryotic transcription factor sequence specificity. <i>Cell</i> , <b>2014</b> , 158, 1431-1443	56.2	866
13	Arabidopsis basic helix-loop-helix transcription factors MYC2, MYC3, and MYC4 regulate glucosinolate biosynthesis, insect performance, and feeding behavior. <i>Plant Cell</i> , <b>2013</b> , 25, 3117-32	11.6	313
12	A trio of viral proteins tunes aphid-plant interactions in Arabidopsis thaliana. <i>PLoS ONE</i> , <b>2013</b> , 8, e8306	63.7	49
11	Transgenerational epigenetic instability is a source of novel methylation variants. <i>Science</i> , <b>2011</b> , 334, 369-73	33.3	485

10	An antiviral defense role of AGO2 in plants. PLoS ONE, <b>2011</b> , 6, e14639	3.7	238
9	Cucumber mosaic virus and its 2b RNA silencing suppressor modify plant-aphid interactions in tobacco. <i>Scientific Reports</i> , <b>2011</b> , 1, 187	4.9	98
8	Symptom induction and RNA silencing suppression by the cucumber mosaic virus 2b protein. <i>Plant Signaling and Behavior</i> , <b>2010</b> , 5, 705-8	2.5	22
7	Signaling in induced resistance. <i>Advances in Virus Research</i> , <b>2010</b> , 76, 57-121	10.7	116
6	Cucumber mosaic virus 2b protein subcellular targets and interactions: their significance to RNA silencing suppressor activity. <i>Molecular Plant-Microbe Interactions</i> , <b>2010</b> , 23, 294-303	3.6	147
5	Disruption of two defensive signaling pathways by a viral RNA silencing suppressor. <i>Molecular Plant-Microbe Interactions</i> , <b>2010</b> , 23, 835-45	3.6	144
4	RNA silencing in plants: Flash report!. Silence: A Journal of RNA Regulation, 2010, 1, 13		7
3	Effects of DICER-like proteins 2, 3 and 4 on cucumber mosaic virus and tobacco mosaic virus infections in salicylic acid-treated plants. <i>Journal of General Virology</i> , <b>2009</b> , 90, 3010-3014	4.9	41
2	The role of the Cucumber mosaic virus 2b protein in viral movement and symptom induction. <i>Molecular Plant-Microbe Interactions</i> , <b>2009</b> , 22, 642-54	3.6	92
1	Selective targeting of miRNA-regulated plant development by a viral counter-silencing protein.  Plant Journal, 2007, 50, 240-52	6.9	106