## Matthew M S Evans

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Vision, challenges and opportunities for a Plant Cell Atlas. ELife, 2021, 10, .	6.0	31
2	Insights into the molecular control of cross-incompatibility in Zea mays. Plant Reproduction, 2020, 33, 117-128.	2.2	21
3	High expression in maize pollen correlates with genetic contributions to pollen fitness as well as with coordinated transcription from neighboring transposable elements. PLoS Genetics, 2020, 16, e1008462.	3.5	30
4	Maternal regulation of seed growth and patterning in flowering plants. Current Topics in Developmental Biology, 2020, 140, 257-282.	2.2	10
5	Title is missing!. , 2020, 16, e1008462.		0
6	Title is missing!. , 2020, 16, e1008462.		0
7	Title is missing!. , 2020, 16, e1008462.		0
8	Title is missing!. , 2020, 16, e1008462.		0
9	Title is missing!. , 2020, 16, e1008462.		0
10	A pistil-expressed pectin methylesterase confers cross-incompatibility between strains of Zea mays. Nature Communications, 2019, 10, 2304.	12.8	26
11	RNA Isolation and Analysis of LncRNAs from Gametophytes of Maize. Methods in Molecular Biology, 2019, 1933, 67-86.	0.9	3
12	Live-Cell Imaging of Auxin and Cytokinin Signaling in Maize Female Gametophytes. Methods in Molecular Biology, 2017, 1669, 95-101.	0.9	3
13	Maternal Gametophyte Effects on Seed Development in Maize. Genetics, 2016, 204, 233-248.	2.9	17
14	Parent-of-Origin-Effect <i>rough endosperm</i> Mutants in Maize. Genetics, 2016, 204, 221-231.	2.9	16
15	Correlation between a loss of auxin signaling and a loss of proliferation in maize antipodal cells. Frontiers in Plant Science, 2015, 6, 187.	3.6	31
16	Discovery of novel transcripts and gametophytic functions via RNA-seq analysis of maize gametophytic transcriptomes. Genome Biology, 2014, 15, 414.	8.8	74
17	Genome-wide discovery and characterization of maize long non-coding RNAs. Genome Biology, 2014, 15, R40.	9.6	419
18	Genetic and cellular analysis of cross-incompatibility in Zea mays. Plant Reproduction, 2014, 27, 19-29.	2.2	30

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19	Analysis of <i>stunter1</i> , a Maize Mutant with Reduced Gametophyte Size and Maternal Effects on Seed Development. Genetics, 2011, 187, 1085-1097.	2.9	20
20	The Zea mays Sexual Compatibility Gene ga2: Naturally Occurring Alleles, Their Distribution, and Role in Reproductive Isolation. Journal of Heredity, 2010, 101, 737-749.	2.4	50
21	The Maize Megagametophyte. , 2009, , 79-104.		15
22	The indeterminate gametophyte1 Gene of Maize Encodes a LOB Domain Protein Required for Embryo Sac and Leaf Development. Plant Cell, 2007, 19, 46-62.	6.6	211
23	Maternal Gametophytic baseless1 Is Required for Development of the Central Cell and Early Endosperm Patterning in Maize (Zea mays). Genetics, 2006, 174, 317-329.	2.9	54
24	Pollen–pistil barriers to crossing in maize and teosinte result from incongruity rather than active rejection. Sexual Plant Reproduction, 2005, 18, 187-194.	2.2	36
25	Unique features of the plant life cycle and their consequences. Nature Reviews Genetics, 2003, 4, 369-379.	16.3	158
26	Interaction Between Maternal Effect and Zygotic Effect Mutations During Maize Seed Development. Genetics, 2001, 159, 303-315.	2.9	55
27	The viviparous8 mutation delays vegetative phase change and accelerates the rate of seedling growth in maize. Plant Journal, 1997, 12, 769-779.	5.7	31