

# Simon J Conn

## List of Publications by Year in descending order

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Version: 2024-02-01

36  
papers

5,847  
citations

218381

26  
h-index

344852

36  
g-index

37  
all docs

37  
docs citations

37  
times ranked

8783  
citing authors

#	ARTICLE	IF	CITATIONS
1	The RNA Binding Protein Quaking Regulates Formation of circRNAs. <i>Cell</i> , 2015, 160, 1125-1134.	13.5	1,698
2	Wheat grain yield on saline soils is improved by an ancestral Na <sup>+</sup> transporter gene. <i>Nature Biotechnology</i> , 2012, 30, 360-364.	9.4	690
3	A circRNA from SEPALLATA3 regulates splicing of its cognate mRNA through R-loop formation. <i>Nature Plants</i> , 2017, 3, 17053.	4.7	434
4	Pluripotent cell division cycles are driven by ectopic Cdk2, cyclin A/E and E2F activities. <i>Oncogene</i> , 2002, 21, 8320-8333.	2.6	332
5	Comparative physiology of elemental distributions in plants. <i>Annals of Botany</i> , 2010, 105, 1081-1102.	1.4	288
6	Cell-Specific Vacuolar Calcium Storage Mediated by <i>CAX1</i> Regulates Apoplastic Calcium Concentration, Gas Exchange, and Plant Productivity in <i>Arabidopsis</i> . <i>Plant Cell</i> , 2011, 23, 240-257.	3.1	222
7	Calcium delivery and storage in plant leaves: exploring the link with water flow. <i>Journal of Experimental Botany</i> , 2011, 62, 2233-2250.	2.4	208
8	RNA Clamping by Vasa Assembles a piRNA Amplifier Complex on Transposon Transcripts. <i>Cell</i> , 2014, 157, 1698-1711.	13.5	208
9	Xylem ionic relations and salinity tolerance in barley. <i>Plant Journal</i> , 2010, 61, 839-853.	2.8	198
10	Purification, molecular cloning, and characterization of glutathione S-transferases (GSTs) from pigmented <i>Vitis vinifera</i> L. cell suspension cultures as putative anthocyanin transport proteins. <i>Journal of Experimental Botany</i> , 2008, 59, 3621-3634.	2.4	193
11	Protocol: optimising hydroponic growth systems for nutritional and physiological analysis of <i>Arabidopsis thaliana</i> and other plants. <i>Plant Methods</i> , 2013, 9, 4.	1.9	167
12	Developmental Activation of the Rb-E2F Pathway and Establishment of Cell Cycle-regulated Cyclin-dependent Kinase Activity during Embryonic Stem Cell Differentiation. <i>Molecular Biology of the Cell</i> , 2005, 16, 2018-2027.	0.9	152
13	An update on magnesium homeostasis mechanisms in plants. <i>Metallomics</i> , 2013, 5, 1170.	1.0	133
14	The response of the maize nitrate transport system to nitrogen demand and supply across the lifecycle. <i>New Phytologist</i> , 2013, 198, 82-94.	3.5	108
15	Magnesium transporters, MGT2/MRS2 <sup>1</sup> and MGT3/MRS2 <sup>5</sup> , are important for magnesium partitioning within <i>Arabidopsis thaliana</i> mesophyll vacuoles. <i>New Phytologist</i> , 2011, 190, 583-594.	3.5	99
16	Structural Basis for the Oligomerization of the MADS Domain Transcription Factor SEPALLATA3 in <i>Arabidopsis</i> . <i>Plant Cell</i> , 2014, 26, 3603-3615.	3.1	97
17	miR-200/375 control epithelial plasticity-associated alternative splicing by repressing the <i>scp</i> binding protein Quaking. <i>EMBO Journal</i> , 2018, 37, .	3.5	82
18	Tetramerization of MADS family transcription factors SEPALLATA3 and AGAMOUS is required for floral meristem determinacy in <i>Arabidopsis</i> . <i>Nucleic Acids Research</i> , 2018, 46, 4966-4977.	6.5	81

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19	Anthocyanic vacuolar inclusions (AVIs) selectively bind acylated anthocyanins in <i>Vitis vinifera</i> L. (grapevine) suspension culture. <i>Biotechnology Letters</i> , 2003, 25, 835-839.	1.1	62
20	Characterization of anthocyanic vacuolar inclusions in <i>Vitis vinifera</i> L. cell suspension cultures. <i>Planta</i> , 2010, 231, 1343-1360.	1.6	55
21	Protocol: a fast and simple in situ PCR method for localising gene expression in plant tissue. <i>Plant Methods</i> , 2014, 10, 29.	1.9	45
22	Variation for N Uptake System in Maize: Genotypic Response to N Supply. <i>Frontiers in Plant Science</i> , 2015, 6, 936.	1.7	39
23	Heterodimerization of Arabidopsis calcium/proton exchangers contributes to regulation of guard cell dynamics and plant defense responses. <i>Journal of Experimental Botany</i> , 2017, 68, 4171-4183.	2.4	39
24	CircRNAs in Plants. <i>Advances in Experimental Medicine and Biology</i> , 2018, 1087, 329-343.	0.8	37
25	Cell-specific compartmentation of mineral nutrients is an essential mechanism for optimal plant productivity—another role for <i>TPC1</i> ? <i>Plant Signaling and Behavior</i> , 2011, 6, 1656-1661.	1.2	34
26	To Stretch the Boundary of Secondary Metabolite Production in Plant Cell-Based Bioprocessing: Anthocyanin as a Case Study. <i>Journal of Biomedicine and Biotechnology</i> , 2004, 2004, 264-271.	3.0	29
27	SplintQuant: a method for accurately quantifying circular RNA transcript abundance without reverse transcription bias. <i>Rna</i> , 2019, 25, 1202-1210.	1.6	29
28	Exploiting natural variation to uncover candidate genes that control element accumulation in <i>Arabidopsis thaliana</i> . <i>New Phytologist</i> , 2012, 193, 859-866.	3.5	24
29	A Highly Efficient Strategy for Overexpressing circRNAs. <i>Methods in Molecular Biology</i> , 2018, 1724, 97-105.	0.4	16
30	A Neuroethics Framework for the Australian Brain Initiative. <i>Neuron</i> , 2019, 101, 365-369.	3.8	11
31	The Suitability of Glioblastoma Cell Lines as Models for Primary Glioblastoma Cell Metabolism. <i>Cancers</i> , 2020, 12, 3722.	1.7	10
32	Don't go in circles: confounding factors in gene expression profiling. <i>EMBO Journal</i> , 2018, 37, .	3.5	8
33	SRRM4 Expands the Repertoire of Circular RNAs by Regulating Microexon Inclusion. <i>Cells</i> , 2020, 9, 2488.	1.8	8
34	The Non-Coding RNA Journal Club: Highlights on Recent Papers—7. <i>Non-coding RNA</i> , 2019, 5, 40.	1.3	2
35	Transcriptomics on Small Samples. <i>Methods in Molecular Biology</i> , 2012, 913, 335-350.	0.4	2
36	The Non-Coding RNA Journal Club: Highlights on Recent Papers—6. <i>Non-coding RNA</i> , 2018, 4, 23.	1.3	0