

Sylvain Aubry

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6031592/publications.pdf>

Version: 2024-02-01

32
papers

2,724
citations

304743

22
h-index

414414

32
g-index

37
all docs

37
docs citations

37
times ranked

3008
citing authors

#	ARTICLE	IF	CITATIONS
1	Pheophytin Pheophorbide Hydrolase (Pheophytinase) Is Involved in Chlorophyll Breakdown during Leaf Senescence in <i>Arabidopsis</i> Å. <i>Plant Cell</i> , 2009, 21, 767-785.	6.6	513
2	Chlorophyll Breakdown in Senescent <i>Arabidopsis</i> Leaves. Characterization of Chlorophyll Catabolites and of Chlorophyll Catabolic Enzymes Involved in the Degreening Reaction. <i>Plant Physiology</i> , 2005, 139, 52-63.	4.8	278
3	In Vivo Participation of Red Chlorophyll Catabolite Reductase in Chlorophyll Breakdown. <i>Plant Cell</i> , 2007, 19, 369-387.	6.6	215
4	Cross-Species Identification of Mendel's I Locus. <i>Science</i> , 2007, 315, 73-73.	12.6	168
5	The role of proteins in C3 plants prior to their recruitment into the C4 pathway. <i>Journal of Experimental Botany</i> , 2011, 62, 3049-3059.	4.8	168
6	Deep Evolutionary Comparison of Gene Expression Identifies Parallel Recruitment of Trans-Factors in Two Independent Origins of C4 Photosynthesis. <i>PLoS Genetics</i> , 2014, 10, e1004365.	3.5	165
7	From crop to model to crop: identifying the genetic basis of the staygreen mutation in the <i>Lolium</i> / <i>Festuca</i> forage and amenity grasses. <i>New Phytologist</i> , 2006, 172, 592-597.	7.3	98
8	Evolution of GOLDEN2-LIKE gene function in C3 and C4 plants. <i>Planta</i> , 2013, 237, 481-495.	3.2	98
9	Stay-green protein, defective in Mendel's green cotyledon mutant, acts independent and upstream of pheophorbide a oxygenase in the chlorophyll catabolic pathway. <i>Plant Molecular Biology</i> , 2008, 67, 243-256.	3.9	96
10	<i>Arabidopsis</i> uses two gluconeogenic gateways for organic acids to fuel seedling establishment. <i>Nature Communications</i> , 2015, 6, 6659.	12.8	95
11	Accumulation of chlorophyll catabolites photosensitizes the hypersensitive response elicited by <i>Pseudomonas syringae</i> in <i>Arabidopsis</i> . <i>New Phytologist</i> , 2010, 188, 161-174.	7.3	91
12	MES16, a Member of the Methyltransferase Protein Family, Specifically Demethylates Fluorescent Chlorophyll Catabolites during Chlorophyll Breakdown in <i>Arabidopsis</i> . <i>Plant Physiology</i> , 2012, 158, 628-641.	4.8	83
13	Transcript residency on ribosomes reveals a key role for the <i>Arabidopsis thaliana</i> bundle sheath in sulfur and glucosinolate metabolism. <i>Plant Journal</i> , 2014, 78, 659-673.	5.7	83
14	A Role for TIC55 as a Hydroxylase of Phyllobilins, the Products of Chlorophyll Breakdown during Plant Senescence. <i>Plant Cell</i> , 2016, 28, 2510-2527.	6.6	75
15	Molecular evolution of genes recruited into C4 photosynthesis. <i>Trends in Plant Science</i> , 2012, 17, 213-220.	8.8	73
16	An Untranslated cis-Element Regulates the Accumulation of Multiple C ₄ Enzymes in <i>Gynandropsis gynandra</i> Mesophyll Cells. <i>Plant Cell</i> , 2016, 28, 454-465.	6.6	73
17	Circadian oscillations of cytosolic free calcium regulate the <i>Arabidopsis</i> circadian clock. <i>Nature Plants</i> , 2018, 4, 690-698.	9.3	65
18	Contribution of Untargeted Metabolomics for Future Assessment of Biotech Crops. <i>Trends in Plant Science</i> , 2018, 23, 1047-1056.	8.8	45

#	ARTICLE	IF	CITATIONS
19	The Future of Digital Sequence Information for Plant Genetic Resources for Food and Agriculture. <i>Frontiers in Plant Science</i> , 2019, 10, 1046.	3.6	38
20	Non-specific activities of the major herbicide-resistance gene BAR. <i>Nature Plants</i> , 2017, 3, 937-945.	9.3	33
21	Pheophorbide <i>ide</i> May Regulate Jasmonate Signaling during Dark-Induced Senescence. <i>Plant Physiology</i> , 2020, 182, 776-791.	4.8	32
22	A Specific Transcriptome Signature for Guard Cells from the C ₄ Plant <i>Gynandropsis gynandra</i> . <i>Plant Physiology</i> , 2016, 170, 1345-1357.	4.8	29
23	Strigolactones Play an Important Role in Shaping Exodermal Morphology via a KAI2-Dependent Pathway. <i>IScience</i> , 2019, 17, 144-154.	4.1	24
24	The pyruvate, orthophosphate dikinase regulatory proteins of Arabidopsis are both bifunctional and interact with the catalytic and nucleotide-binding domains of pyruvate, orthophosphate dikinase. <i>Plant Journal</i> , 2011, 68, 1070-1080.	5.7	20
25	Post-transcriptional regulation of photosynthetic genes is a key driver of C ₄ leaf ontogeny. <i>Journal of Experimental Botany</i> , 2017, 68, 137-146.	4.8	16
26	Bringing access and benefit sharing into the digital age. <i>Plants People Planet</i> , 2022, 4, 5-12.	3.3	11
27	Evolution of chlorophyll degradation is associated with plant transition to land. <i>Plant Journal</i> , 2022, 109, 1473-1488.	5.7	10
28	Endoreduplication is not involved in bundle-sheath formation in the C ₄ species <i>Cleome gynandra</i> . <i>Journal of Experimental Botany</i> , 2014, 65, 3557-3566.	4.8	7
29	Genetically modified crops in Switzerland: implications for agrosystem sustainability evidenced by multi-criteria model. <i>Agronomy for Sustainable Development</i> , 2016, 36, 1.	5.3	7
30	An evergreen mind and a heart for the colors of fall. <i>Journal of Experimental Botany</i> , 2021, 72, 4625-4633.	4.8	4
31	De novo transcriptome assembly data of the marine bioluminescent dinoflagellate <i>Pyrocystis lunula</i> . <i>Data in Brief</i> , 2021, 37, 107254.	1.0	3
32	Chapter 10. Omics-based Detection, Identification and Quantification of GM Food and Feed: Current Challenges and Perspectives. <i>Food Chemistry, Function and Analysis</i> , 2021, , 257-270.	0.2	1