## Samuel P Hazen

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/1979510/publications.pdf Version: 2024-02-01



SAMILEL D HAZEN

#	Article	IF	CITATIONS
1	Network Discovery Pipeline Elucidates Conserved Time-of-Day–Specific cis-Regulatory Modules. PLoS Genetics, 2008, 4, e14.	3.5	474
2	Brachypodium as a Model for the Grasses: Today and the Future Â. Plant Physiology, 2011, 157, 3-13.	4.8	243
3	A Morning-Specific Phytohormone Gene Expression Program underlying Rhythmic Plant Growth. PLoS Biology, 2008, 6, e225.	5.6	197
4	Cellulose Synthase-Like Genes of Rice. Plant Physiology, 2002, 128, 336-340.	4.8	178
5	Expression profiling of rice segregating for drought tolerance QTLs using a rice genome array. Functional and Integrative Genomics, 2005, 5, 104-116.	3.5	103
6	Exploring the transcriptional landscape of plant circadian rhythms using genome tiling arrays. Genome Biology, 2009, 10, R17.	9.6	103
7	A High-Resolution Map of Arabidopsis Recombinant Inbred Lines by Whole-Genome Exon Array Hybridization. PLoS Genetics, 2006, 2, e144.	3.5	97
8	Rapid Array Mapping of Circadian Clock and Developmental Mutations in Arabidopsis. Plant Physiology, 2005, 138, 990-997.	4.8	85
9	Gene expression profiling of plant responses to abiotic stress. Functional and Integrative Genomics, 2003, 3, 105-111.	3.5	84
10	Functional characterization of cinnamyl alcohol dehydrogenase and caffeic acid O-methyltransferase in Brachypodium distachyon. BMC Biotechnology, 2013, 13, 61.	3.3	84
11	Perturbation ofBrachypodium distachyon CELLULOSE SYNTHASE A4or7results in abnormal cell walls. BMC Plant Biology, 2013, 13, 131.	3.6	81
12	Environmental niche variation and evolutionary diversification of the <i>Brachypodium distachyon</i> grass complex species in their native circumâ€Mediterranean range. American Journal of Botany, 2015, 102, 1073-1088.	1.7	73
13	Gradual polyploid genome evolution revealed by pan-genomic analysis of Brachypodium hybridum and its diploid progenitors. Nature Communications, 2020, 11, 3670.	12.8	67
14	Quantitative Trait Loci and Comparative Genomics of Cereal Cell Wall Composition. Plant Physiology, 2003, 132, 263-271.	4.8	64
15	Transcriptional Regulation of Grass Secondary Cell Wall Biosynthesis: Playing Catch-Up with Arabidopsis thaliana. Frontiers in Plant Science, 2012, 3, 74.	3.6	61
16	Genotype, development and tissue-derived variation of cell-wall properties in the lignocellulosic energy crop Miscanthus. Annals of Botany, 2014, 114, 1265-1277.	2.9	56
17	A stressâ€associated protein, AtSAP13, from <scp><i>Arabidopsis thaliana</i></scp> provides tolerance to multiple abiotic stresses. Plant, Cell and Environment, 2018, 41, 1171-1185.	5.7	52
18	Climate-smart crops with enhanced photosynthesis. Journal of Experimental Botany, 2018, 69, 3801-3809.	4.8	50

SAMUEL P HAZEN

#	Article	IF	CITATIONS
19	Gene arrays are not just for measuring gene expression. Trends in Plant Science, 2003, 8, 413-416.	8.8	47
20	Daily Changes in Temperature, Not the Circadian Clock, Regulate Growth Rate in Brachypodium distachyon. PLoS ONE, 2014, 9, e100072.	2.5	47
21	A cell wall reference profile for <i>Miscanthus</i> bioenergy crops highlights compositional and structural variations associated withÂdevelopment and organ origin. New Phytologist, 2017, 213, 1710-1725.	7.3	44
22	Grass secondary cell walls, <i>Brachypodium distachyon</i> as a model for discovery. New Phytologist, 2020, 227, 1649-1667.	7.3	40
23	Regulation of Cell Wall Thickening by a Medley of Mechanisms. Trends in Plant Science, 2019, 24, 853-866.	8.8	34
24	Cell Walls and the Developmental Anatomy of the Brachypodium distachyon Stem Internode. PLoS ONE, 2013, 8, e80640.	2.5	34
25	AFLP in Triticum aestivum L.: patterns of genetic diversity and genome distribution. Euphytica, 2002, 125, 89-102.	1.2	29
26	Biological conversion assay using Clostridium phytofermentans to estimate plant feedstock quality. Biotechnology for Biofuels, 2012, 5, 5.	6.2	28
27	PIL1 Participates in a Negative Feedback Loop that Regulates Its Own Gene Expression in Response to Shade. Molecular Plant, 2014, 7, 1582-1585.	8.3	27
28	Lignocellulosic feedstocks: research progress and challenges in optimizing biomass quality and yield. Frontiers in Plant Science, 2013, 4, 474.	3.6	21
29	Brachypodium: 20 years as a grass biology model system; the way forward?. Trends in Plant Science, 2022, 27, 1002-1016.	8.8	21
30	<i><scp>SECONDARY WALL ASSOCIATED MYB</scp>1</i> is a positive regulator of secondary cell wall thickening in <i>Brachypodium distachyon</i> and is not found in the Brassicaceae. Plant Journal, 2018, 96, 532-545.	5.7	20
31	Changes in ambient temperature are the prevailing cue in determining <i>Brachypodium distachyon</i> diurnal gene regulation. New Phytologist, 2020, 227, 1709-1724.	7.3	16
32	Rice Genome-Scale Network Integration Reveals Transcriptional Regulators of Grass Cell Wall Synthesis. Frontiers in Plant Science, 2019, 10, 1275.	3.6	14
33	Cellulose Synthase-Like Genes of Rice. Plant Physiology, 2002, 128, 336-340.	4.8	14
34	Secondary Wall Regulating NACs Differentially Bind at the Promoter at a CELLULOSE SYNTHASE A4 Cis-eQTL. Frontiers in Plant Science, 2018, 9, 1895.	3.6	11
35	Title is missing!. Genetic Resources and Crop Evolution, 2002, 49, 439-448.	1.6	9
36	A High-Throughput Biological Conversion Assay for Determining Lignocellulosic Quality. Methods in Molecular Biology, 2012, 918, 341-349.	0.9	5

#	Article	IF	CITATIONS
37	Direct Image-Based Enumeration of Clostridium phytofermentans Cells on Insoluble Plant Biomass Growth Substrates. Applied and Environmental Microbiology, 2016, 82, 972-978.	3.1	1