

Ove Nilsson

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2889742/publications.pdf>

Version: 2024-02-01

51
papers

6,349
citations

159585

30
h-index

206112

48
g-index

55
all docs

55
docs citations

55
times ranked

7304
citing authors

#	ARTICLE	IF	CITATIONS
1	The Norway spruce genome sequence and conifer genome evolution. <i>Nature</i> , 2013, 497, 579-584.	27.8	1,303
2	CO/FT Regulatory Module Controls Timing of Flowering and Seasonal Growth Cessation in Trees. <i>Science</i> , 2006, 312, 1040-1043.	12.6	904
3	A developmental switch sufficient for flower initiation in diverse plants. <i>Nature</i> , 1995, 377, 495-500.	27.8	787
4	Gibberellins Promote Flowering of Arabidopsis by Activating the LEAFY Promoter. <i>Plant Cell</i> , 1998, 10, 791-800.	6.6	519
5	A transcriptional timetable of autumn senescence. <i>Genome Biology</i> , 2004, 5, R24.	9.6	226
6	AspWood: High-Spatial-Resolution Transcriptome Profiles Reveal Uncharacterized Modularity of Wood Formation in <i>Populus tremula</i> . <i>Plant Cell</i> , 2017, 29, 1585-1604.	6.6	219
7	The BLADE ON PETIOLE genes act redundantly to control the growth and development of lateral organs. <i>Development (Cambridge)</i> , 2005, 132, 2203-2213.	2.5	207
8	Electronic plants. <i>Science Advances</i> , 2015, 1, e1501136.	10.3	190
9	Insights into Conifer Giga-Genomes. <i>Plant Physiology</i> , 2014, 166, 1724-1732.	4.8	164
10	Flowering-Time Genes Modulate the Response to LEAFY Activity. <i>Genetics</i> , 1998, 150, 403-410.	2.9	151
11	Spatial pattern of cauliflower mosaic virus 35S promoter-luciferase expression in transgenic hybrid aspen trees monitored by enzymatic assay and non-destructive imaging. <i>Transgenic Research</i> , 1992, 1, 209-220.	2.4	138
12	<i>WUSCHEL</i> -RELATED HOMEODOMAIN 4 (<i>WOX4</i>)-like genes regulate cambial cell division activity and secondary growth in <i>Populus</i> trees. <i>New Phytologist</i> , 2017, 215, 642-657.	7.3	117
13	BLADE-ON-PETIOLE proteins act in an E3 ubiquitin ligase complex to regulate PHYTOCHROME INTERACTING FACTOR 4 abundance. <i>Elife</i> , 2017, 6, .	6.0	106
14	Class I KNOX transcription factors promote differentiation of cambial derivatives into xylem fibers in the <i>Arabidopsis</i> hypocotyl. <i>Development (Cambridge)</i> , 2014, 141, 4311-4319.	2.5	97
15	Analysis of conifer <i>FLOWERING LOCUS T</i> / <i>TERMINAL FLOWER1</i> -like genes provides evidence for dramatic biochemical evolution in the angiosperm <i>FT</i> lineage. <i>New Phytologist</i> , 2012, 196, 1260-1273.	7.3	90
16	Indole-3-acetic acid homeostasis in transgenic tobacco plants expressing the <i>Agrobacterium rhizogenes</i> rolB gene. <i>Plant Journal</i> , 1993, 3, 681-689.	5.7	89
17	Revisiting tree maturation and floral initiation in the poplar functional genomics era. <i>New Phytologist</i> , 2004, 164, 43-51.	7.3	88
18	The Arabidopsis LRR-RLK, PXC1, is a regulator of secondary wall formation correlated with the TDIF-PXY/TDR-WOX4 signaling pathway. <i>BMC Plant Biology</i> , 2013, 13, 94.	3.6	80

#	ARTICLE	IF	CITATIONS
19	A major locus controls local adaptation and adaptive life history variation in a perennial plant. <i>Genome Biology</i> , 2018, 19, 72.	8.8	76
20	NorWood: a gene expression resource for evo–devo studies of conifer wood development. <i>New Phytologist</i> , 2017, 216, 482-494.	7.3	71
21	<i>FT</i> overexpression induces precocious flowering and normal reproductive development in <i>Eucalyptus</i>. <i>Plant Biotechnology Journal</i> , 2016, 14, 808-819.	8.3	70
22	Molecular regulation of phenology in trees â€” because the seasons they are a-changinâ€™. <i>Current Opinion in Plant Biology</i> , 2016, 29, 73-79.	7.1	70
23	Expression of two heterologous promoters, <i>Agrobacterium rhizogenes</i> rolC and cauliflower mosaic virus 35S, in the stem of transgenic hybrid aspen plants during the annual cycle of growth and dormancy. <i>Plant Molecular Biology</i> , 1996, 31, 887-895.	3.9	57
24	<i>GIGANTEA</i>-like genes control seasonal growth cessation in <i>Populus</i>. <i>New Phytologist</i> , 2018, 218, 1491-1503.	7.3	55
25	Transcriptional Roadmap to Seasonal Variation in Wood Formation of Norway Spruce. <i>Plant Physiology</i> , 2018, 176, 2851-2870.	4.8	40
26	Autumn senescence in aspen is not triggered by day length. <i>Physiologia Plantarum</i> , 2018, 162, 123-134.	5.2	40
27	Modulating the timing of flowering. <i>Current Opinion in Biotechnology</i> , 1997, 8, 195-199.	6.6	39
28	The <i>Agrobacterium rhizogenes</i> rolB and rolC promoters are expressed in pericycle cells competent to serve as root initials in transgenic hybrid aspen. <i>Physiologia Plantarum</i> , 1997, 100, 456-462.	5.2	35
29	Getting to the root: The role of the <i>Agrobacterium rhizogenes</i> rol genes in the formation of hairy roots. <i>Physiologia Plantarum</i> , 1997, 100, 463-473.	5.2	35
30	<sc>CLE</sc> peptide signaling in plantsâ€™â€”the power of moving around. <i>Physiologia Plantarum</i> , 2015, 155, 74-87.	5.2	33
31	<sc>LEAFY</sc> activity is post–transcriptionally regulated by <sc>BLADE ON PETIOLE</sc>2 and <sc>CULLIN</sc>3 in <i>Arabidopsis</i> . <i>New Phytologist</i> , 2018, 220, 579-592.	7.3	32
32	Phytochrome B and PHYTOCHROME INTERACTING FACTOR8 modulate seasonal growth in trees. <i>New Phytologist</i> , 2021, 232, 2339-2352.	7.3	31
33	FLOWERING LOCUS T paralogs control the annual growth cycle in <i>Populus</i> trees. <i>Current Biology</i> , 2022, 32, 2988-2996.e4.	3.9	24
34	Successful crossings with early flowering transgenic poplar: interspecific crossings, but not transgenesis, promoted aberrant phenotypes in offspring. <i>Plant Biotechnology Journal</i> , 2014, 12, 1066-1074.	8.3	20
35	Low temperatures are required to induce the development of fertile flowers in transgenic male and female early flowering poplar (<i>Populus tremula</i> L.). <i>Tree Physiology</i> , 2016, 36, 667-677.	3.1	19
36	Transcriptome analysis of embryonic domains in Norway spruce reveals potential regulators of suspensor cell death. <i>PLoS ONE</i> , 2018, 13, e0192945.	2.5	17

#	ARTICLE	IF	CITATIONS
37	Separation and identification of cytokinins using combined capillary liquid chromatography/mass spectrometry. <i>Biological Mass Spectrometry</i> , 1993, 22, 201-210.	0.5	14
38	Peptide encoding <i>Populus CLV3/ESR</i> -RELATED 47 (<i>PttCLE47</i>) promotes cambial development and secondary xylem formation in hybrid aspen. <i>New Phytologist</i> , 2020, 226, 75-85.	7.3	13
39	Functional metabolomics as a tool to analyze Mediator function and structure in plants. <i>PLoS ONE</i> , 2017, 12, e0179640.	2.5	13
40	Novel monomeric luciferase enzymes as tools to study plant gene regulation <i>in vivo</i> . <i>Luminescence</i> , 1990, 5, 79-87.	0.0	12
41	Certification for gene-edited forests. <i>Science</i> , 2019, 365, 767-768.	12.6	12
42	Integrative Analysis of Three RNA Sequencing Methods Identifies Mutually Exclusive Exons of MADS-Box Isoforms During Early Bud Development in <i>Picea abies</i> . <i>Frontiers in Plant Science</i> , 2018, 9, 1625.	3.6	10
43	<i>Populus SVL</i> Acts in Leaves to Modulate the Timing of Growth Cessation and Bud Set. <i>Frontiers in Plant Science</i> , 2022, 13, 823019.	3.6	8
44	EU Regulations Impede Market Introduction of GM Forest Trees. <i>Trends in Plant Science</i> , 2016, 21, 283-285.	8.8	6
45	<i>GIGANTEA</i> influences leaf senescence in trees in two different ways. <i>Plant Physiology</i> , 2021, 187, 2435-2450.	4.8	5
46	Indole-3-acetic acid homeostasis in transgenic tobacco plants expressing the <i>Agrobacterium rhizogenes rolB</i> gene. <i>Plant Journal</i> , 1993, 3, 681-689.	5.7	5
47	Plant Evolution: Measuring the Length of the Day. <i>Current Biology</i> , 2009, 19, R302-R303.	3.9	4
48	The <i>Agrobacterium rhizogenes rolB</i> and <i>rolC</i> promoters are expressed in pericycle cells competent to serve as root initials in transgenic hybrid aspen. <i>Physiologia Plantarum</i> , 1997, 100, 456-462.	5.2	4
49	Photoperiodic Control of Dormancy and Flowering in Trees. , 2009, , 88-106.		1
50	Arabidopsis Research 2000. <i>Plant Cell</i> , 2000, 12, 2302.	6.6	0
51	Variation in non-target traits in genetically modified hybrid aspens does not exceed natural variation. <i>New Biotechnology</i> , 2021, 64, 27-36.	4.4	0