

Zheng-Hua Ye

List of Publications by Year in descending order

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79
papers

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36271

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80
all docs

80
docs citations

80
times ranked

6690
citing authors

#	ARTICLE	IF	CITATIONS
1	Cell wall biology of the moss <i>Physcomitrium patens</i> . Journal of Experimental Botany, 2022, 73, 4440-4453.	2.4	10
2	XND1 Regulates Secondary Wall Deposition in Xylem Vessels through the Inhibition of VND Functions. Plant and Cell Physiology, 2021, 62, 53-65.	1.5	20
3	Xylem vessel-specific SND5 and its homologs regulate secondary wall biosynthesis through activating secondary wall NAC binding elements. New Phytologist, 2021, 231, 1496-1509.	3.5	24
4	A Single Xyloglucan Xylosyltransferase Is Sufficient for Generation of the XXXG Xylosylation Pattern of Xyloglucan. Plant and Cell Physiology, 2021, 62, 1589-1602.	1.5	8
5	Functional analysis of GT61 glycosyltransferases from grass species in xylan substitutions. Planta, 2021, 254, 131.	1.6	7
6	Cytosolic Acetyl-CoA Generated by ATP-Citrate Lyase Is Essential for Acetylation of Cell Wall Polysaccharides. Plant and Cell Physiology, 2020, 61, 64-75.	1.5	11
7	A Group of O-Acetyltransferases Catalyze Xyloglucan Backbone Acetylation and Can Alter Xyloglucan Xylosylation Pattern and Plant Growth When Expressed in Arabidopsis. Plant and Cell Physiology, 2020, 61, 1064-1079.	1.5	14
8	Evolutionary origin of O-acetyltransferases responsible for glucomannan acetylation in land plants. New Phytologist, 2019, 224, 466-479.	3.5	26
9	Secondary cell wall biosynthesis. New Phytologist, 2019, 221, 1703-1723.	3.5	185
10	A Novel Rice Xylosyltransferase Catalyzes the Addition of 2-O-Xylosyl Side Chains onto the Xylan Backbone. Plant and Cell Physiology, 2018, 59, 554-565.	1.5	40
11	Biochemical characterization of rice xylan O-acetyltransferases. Planta, 2018, 247, 1489-1498.	1.6	19
12	A group of Populus trichocarpa DUF231 proteins exhibit differential O-acetyltransferase activities toward xylan. PLoS ONE, 2018, 13, e0194532.	1.1	21
13	Xyloglucan O-acetyltransferases from Arabidopsis thaliana and Populus trichocarpa catalyze acetylation of fucosylated galactose residues on xyloglucan side chains. Planta, 2018, 248, 1159-1171.	1.6	26
14	Members of the DUF231 Family are O-Acetyltransferases Catalyzing 2-O- and 3-O-Acetylation of Mannan. Plant and Cell Physiology, 2018, 59, 2339-2349.	1.5	14
15	Cytosol-Localized UDP-Xylose Synthases Provide the Major Source of UDP-Xylose for the Biosynthesis of Xylan and Xyloglucan. Plant and Cell Physiology, 2017, 58, pcw179.	1.5	25
16	Regiospecific Acetylation of Xylan is Mediated by a Group of DUF231-Containing O-Acetyltransferases. Plant and Cell Physiology, 2017, 58, 2126-2138.	1.5	58
17	Mutations of Arabidopsis TBL32 and TBL33 Affect Xylan Acetylation and Secondary Wall Deposition. PLoS ONE, 2016, 11, e0146460.	1.1	76
18	Evolutionary Conservation of Xylan Biosynthetic Genes in <i>Selaginella moellendorffii</i> and <i>Physcomitrella patens</i> . Plant and Cell Physiology, 2016, 57, 1707-1719.	1.5	16

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19	Roles of Arabidopsis TBL34 and TBL35 in xylan acetylation and plant growth. <i>Plant Science</i> , 2016, 243, 120-130.	1.7	59
20	TBL3 and TBL31, Two Arabidopsis DUF231 Domain Proteins, are Required for 3-O-Monoacetylation of Xylan. <i>Plant and Cell Physiology</i> , 2016, 57, 35-45.	1.5	47
21	Functional Characterization of NAC and MYB Transcription Factors Involved in Regulation of Biomass Production in Switchgrass (<i>Panicum virgatum</i>). <i>PLoS ONE</i> , 2015, 10, e0134611.	1.1	68
22	Kinesin-4 Functions in Vesicular Transport on Cortical Microtubules and Regulates Cell Wall Mechanics during Cell Elongation in Plants. <i>Molecular Plant</i> , 2015, 8, 1011-1023.	3.9	83
23	Molecular control of wood formation in trees. <i>Journal of Experimental Botany</i> , 2015, 66, 4119-4131.	2.4	148
24	The Arabidopsis NAC transcription factor NST2 functions together with SND1 and NST1 to regulate secondary wall biosynthesis in fibers of inflorescence stems. <i>Plant Signaling and Behavior</i> , 2015, 10, e989746.	1.2	53
25	Secondary Cell Walls: Biosynthesis, Patterned Deposition and Transcriptional Regulation. <i>Plant and Cell Physiology</i> , 2015, 56, 195-214.	1.5	360
26	Identification and Biochemical Characterization of Four Wood-Associated Glucuronoxylan Methyltransferases in <i>Populus</i> . <i>PLoS ONE</i> , 2014, 9, e87370.	1.1	17
27	Functional roles of rice glycosyltransferase family GT43 in xylan biosynthesis. <i>Plant Signaling and Behavior</i> , 2014, 9, e27809.	1.2	54
28	Identification of a disaccharide side chain 2-O-β-D-galactopyranosyl-β-D-glucuronic acid in Arabidopsis xylan. <i>Plant Signaling and Behavior</i> , 2014, 9, e27933.	1.2	18
29	Alterations of the degree of xylan acetylation in Arabidopsis xylan mutants. <i>Plant Signaling and Behavior</i> , 2014, 9, e27797.	1.2	17
30	Complexity of the transcriptional network controlling secondary wall biosynthesis. <i>Plant Science</i> , 2014, 229, 193-207.	1.7	124
31	Modification of the degree of 4-O-methylation of secondary wall glucuronoxylan. <i>Plant Science</i> , 2014, 219-220, 42-50.	1.7	28
32	Arabidopsis NAC Domain Proteins, VND1 to VND5, Are Transcriptional Regulators of Secondary Wall Biosynthesis in Vessels. <i>PLoS ONE</i> , 2014, 9, e105726.	1.1	169
33	The Arabidopsis DUF231 Domain-Containing Protein ESK1 Mediates 2-O- and 3-O-Acetylation of Xylosyl Residues in Xylan. <i>Plant and Cell Physiology</i> , 2013, 54, 1186-1199.	1.5	129
34	The Poplar MYB Master Switches Bind to the SMRE Site and Activate the Secondary Wall Biosynthetic Program during Wood Formation. <i>PLoS ONE</i> , 2013, 8, e69219.	1.1	130
35	Arabidopsis Family GT43 Members are Xylan Xylosyltransferases Required for the Elongation of the Xylan Backbone. <i>Plant and Cell Physiology</i> , 2012, 53, 135-143.	1.5	76
36	Biochemical characterization of xylan xylosyltransferases involved in wood formation in poplar. <i>Plant Signaling and Behavior</i> , 2012, 7, 332-337.	1.2	23

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37	Arabidopsis GUX Proteins Are Glucuronyltransferases Responsible for the Addition of Glucuronic Acid Side Chains onto Xylan. <i>Plant and Cell Physiology</i> , 2012, 53, 1204-1216.	1.5	97
38	MYB46 and MYB83 Bind to the SMRE Sites and Directly Activate a Suite of Transcription Factors and Secondary Wall Biosynthetic Genes. <i>Plant and Cell Physiology</i> , 2012, 53, 368-380.	1.5	325
39	Three Arabidopsis DUF579 Domain-Containing GXM Proteins are Methyltransferases Catalyzing 4-O-Methylation of Glucuronic Acid on Xylan. <i>Plant and Cell Physiology</i> , 2012, 53, 1934-1949.	1.5	84
40	Molecular Dissection of Xylan Biosynthesis during Wood Formation in Poplar. <i>Molecular Plant</i> , 2011, 4, 730-747.	3.9	81
41	The Four Arabidopsis REDUCED WALL ACETYLATION Genes are Expressed in Secondary Wall-Containing Cells and Required for the Acetylation of Xylan. <i>Plant and Cell Physiology</i> , 2011, 52, 1289-1301.	1.5	132
42	Dissection of the Transcriptional Program Regulating Secondary Wall Biosynthesis during Wood Formation in Poplar. <i>Plant Physiology</i> , 2011, 157, 1452-1468.	2.3	220
43	Transcriptional Activation of Secondary Wall Biosynthesis by Rice and Maize NAC and MYB Transcription Factors. <i>Plant and Cell Physiology</i> , 2011, 52, 1856-1871.	1.5	270
44	The Arabidopsis Family GT43 Glycosyltransferases Form Two Functionally Nonredundant Groups Essential for the Elongation of Glucuronoxylan Backbone. <i>Plant Physiology</i> , 2010, 153, 526-541.	2.3	99
45	Functional Characterization of Poplar Wood-Associated NAC Domain Transcription Factors. <i>Plant Physiology</i> , 2010, 152, 1044-1055.	2.3	245
46	The Poplar MYB Transcription Factors, PtrMYB3 and PtrMYB20, are Involved in the Regulation of Secondary Wall Biosynthesis. <i>Plant and Cell Physiology</i> , 2010, 51, 1084-1090.	1.5	199
47	The poplar PtrWNDs are transcriptional activators of secondary cell wall biosynthesis. <i>Plant Signaling and Behavior</i> , 2010, 5, 469-472.	1.2	48
48	Evolutionary conservation of the transcriptional network regulating secondary cell wall biosynthesis. <i>Trends in Plant Science</i> , 2010, 15, 625-632.	4.3	288
49	Global Analysis of Direct Targets of Secondary Wall NAC Master Switches in Arabidopsis. <i>Molecular Plant</i> , 2010, 3, 1087-1103.	3.9	323
50	Down-Regulation of PoGT47C Expression in Poplar Results in a Reduced Glucuronoxylan Content and an Increased Wood Digestibility by Cellulase. <i>Plant and Cell Physiology</i> , 2009, 50, 1075-1089.	1.5	111
51	Transcriptional regulation of lignin biosynthesis. <i>Plant Signaling and Behavior</i> , 2009, 4, 1028-1034.	1.2	215
52	The Poplar GT8E and GT8F Glycosyltransferases are Functional Orthologs of Arabidopsis PARVUS Involved in Glucuronoxylan Biosynthesis. <i>Plant and Cell Physiology</i> , 2009, 50, 1982-1987.	1.5	37
53	The F8H Glycosyltransferase is a Functional Paralog of FRA8 Involved in Glucuronoxylan Biosynthesis in Arabidopsis. <i>Plant and Cell Physiology</i> , 2009, 50, 812-827.	1.5	98
54	MYB83 Is a Direct Target of SND1 and Acts Redundantly with MYB46 in the Regulation of Secondary Cell Wall Biosynthesis in Arabidopsis. <i>Plant and Cell Physiology</i> , 2009, 50, 1950-1964.	1.5	471

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55	MYB58 and MYB63 Are Transcriptional Activators of the Lignin Biosynthetic Pathway during Secondary Cell Wall Formation in <i>Arabidopsis</i> . <i>Plant Cell</i> , 2009, 21, 248-266.	3.1	737
56	A Battery of Transcription Factors Involved in the Regulation of Secondary Cell Wall Biosynthesis in <i>Arabidopsis</i> . <i>Plant Cell</i> , 2008, 20, 2763-2782.	3.1	866
57	The MYB46 Transcription Factor Is a Direct Target of SND1 and Regulates Secondary Wall Biosynthesis in <i>Arabidopsis</i> . <i>Plant Cell</i> , 2007, 19, 2776-2792.	3.1	576
58	The PARVUS Gene is Expressed in Cells Undergoing Secondary Wall Thickening and is Essential for Glucuronoxylan Biosynthesis. <i>Plant and Cell Physiology</i> , 2007, 48, 1659-1672.	1.5	161
59	<i>Arabidopsis</i> irregular xylem8 and irregular xylem9: Implications for the Complexity of Glucuronoxylan Biosynthesis. <i>Plant Cell</i> , 2007, 19, 549-563.	3.1	396
60	The irregular xylem9 Mutant is Deficient in Xylan Xylosyltransferase Activity. <i>Plant and Cell Physiology</i> , 2007, 48, 1624-1634.	1.5	147
61	Alteration in Secondary Wall Deposition by Overexpression of the <i>Fragile Fiber1</i> Kinesin-Like Protein in <i>Arabidopsis</i> . <i>Journal of Integrative Plant Biology</i> , 2007, 49, 1235-1243.	4.1	17
62	Two NAC domain transcription factors, SND1 and NST1, function redundantly in regulation of secondary wall synthesis in fibers of <i>Arabidopsis</i> . <i>Planta</i> , 2007, 225, 1603-1611.	1.6	373
63	Important new players in secondary wall synthesis. <i>Trends in Plant Science</i> , 2006, 11, 162-164.	4.3	21
64	Disruption of Cortical Microtubules by Overexpression of Green Fluorescent Protein-Tagged alpha-Tubulin 6 Causes a Marked Reduction in Cell Wall Synthesis. <i>Journal of Integrative Plant Biology</i> , 2006, 48, 85-98.	4.1	47
65	SND1, a NAC Domain Transcription Factor, Is a Key Regulator of Secondary Wall Synthesis in Fibers of <i>Arabidopsis</i> . <i>Plant Cell</i> , 2006, 18, 3158-3170.	3.1	655
66	<i>Arabidopsis</i> Fragile Fiber8, Which Encodes a Putative Glucuronyltransferase, Is Essential for Normal Secondary Wall Synthesis. <i>Plant Cell</i> , 2005, 17, 3390-3408.	3.1	301
67	amphivasal vascular bundle 1, a Gain-of-Function Mutation of the IFL1/REV Gene, Is Associated with Alterations in the Polarity of Leaves, Stems and Carpels. <i>Plant and Cell Physiology</i> , 2004, 45, 369-385.	1.5	186
68	Expression of a Mutant Form of Cellulose Synthase AtCesA7 Causes Dominant Negative Effect on Cellulose Biosynthesis. <i>Plant Physiology</i> , 2003, 132, 786-795.	2.3	128
69	Alteration of Oriented Deposition of Cellulose Microfibrils by Mutation of a Katanin-Like Microtubule-Severing Protein. <i>Plant Cell</i> , 2002, 14, 2145-2160.	3.1	248
70	VASCULARTISSUEDIFFERENTIATION ANDPATTERNFORMATION INPLANTS. <i>Annual Review of Plant Biology</i> , 2002, 53, 183-202.	8.6	268
71	Vascular development in <i>Arabidopsis</i> . <i>International Review of Cytology</i> , 2002, 220, 225-256.	6.2	54
72	A Katanin-like Protein Regulates Normal Cell Wall gBiosynthesis and Cell Elongation. <i>Plant Cell</i> , 2001, 13, 807-827.	3.1	330

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73	Alteration of Auxin Polar Transport in the Arabidopsisifl1 Mutants. <i>Plant Physiology</i> , 2001, 126, 549-563.	2.3	146
74	Fibers. A Model for Studying Cell Differentiation, Cell Elongation, and Cell Wall Biosynthesis. <i>Plant Physiology</i> , 2001, 126, 477-479.	2.3	48
75	Essential Role of Caffeoyl Coenzyme A O-Methyltransferase in Lignin Biosynthesis in Woody Poplar Plants. <i>Plant Physiology</i> , 2000, 124, 563-578.	2.3	240
76	Ectopic Deposition of Lignin in the Pith of Stems of Two Arabidopsis Mutants. <i>Plant Physiology</i> , 2000, 123, 59-70.	2.3	94
77	Transformation of the Collateral Vascular Bundles into Amphivasal Vascular Bundles in an Arabidopsis Mutant1. <i>Plant Physiology</i> , 1999, 120, 53-64.	2.3	59
78	IFL1, a Gene Regulating Interfascicular Fiber Differentiation in Arabidopsis, Encodes a Homeodomain-Leucine Zipper Protein. <i>Plant Cell</i> , 1999, 11, 2139-2152.	3.1	271
79	Dual Methylation Pathways in Lignin Biosynthesis. <i>Plant Cell</i> , 1998, 10, 2033-2045.	3.1	239