

# A Genomic Approach to Suberin Biosynthesis and Cork

Plant Physiology

144, 419-431

DOI: [10.1104/pp.106.094227](https://doi.org/10.1104/pp.106.094227)

Citation Report

#	ARTICLE	IF	CITATIONS
1	Identification of acyltransferases required for cutin biosynthesis and production of cutin with suberin-like monomers. Proceedings of the National Academy of Sciences of the United States of America, 2007, 104, 18339-18344.	3.3	348
2	An ABC Transporter Gene of Arabidopsis thaliana, AtWBC11, is Involved in Cuticle Development and Prevention of Organ Fusion. Plant and Cell Physiology, 2007, 48, 1790-1802.	1.5	149
3	Building lipid barriers: biosynthesis of cutin and suberin. Trends in Plant Science, 2008, 13, 236-246.	4.3	779
4	Differential gene expression induced by growth hormone treatment in the uremic rat growth plate. Growth Hormone and IGF Research, 2008, 18, 353-359.	0.5	4
5	Seasonal variation in transcript abundance in cork tissue analyzed by real time RT-PCR. Tree Physiology, 2008, 28, 743-751.	1.4	43
6	The Arabidopsis cytochrome P450 CYP86A1 encodes a fatty acid ̢-hydroxylase involved in suberin monomer biosynthesis. Journal of Experimental Botany, 2008, 59, 2347-2360.	2.4	238
7	CYP86B1 Is Required for Very Long Chain ̢-Hydroxyacid and ̢-Dicarboxylic Acid Synthesis in Root and Seed Suberin Polyester ̂̂. Plant Physiology, 2009, 150, 1831-1843.	2.3	173
8	CYP86A33-Targeted Gene Silencing in Potato Tuber Alters Suberin Composition, Distorts Suberin Lamellae, and Impairs the Periderm's Water Barrier Function ̂̂̂. Plant Physiology, 2009, 149, 1050-1060.	2.3	120
9	Silencing of StKCS6 in potato periderm leads to reduced chain lengths of suberin and wax compounds and increased peridermal transpiration. Journal of Experimental Botany, 2009, 60, 697-707.	2.4	95
10	Identification of an Arabidopsis Feruloyl-Coenzyme A Transferase Required for Suberin Synthesis ̂̂. Plant Physiology, 2009, 151, 1317-1328.	2.3	193
11	Proteomic analysis from haploid and diploid embryos of Quercus suber L. identifies qualitative and quantitative differential expression patterns. Proteomics, 2009, 9, 4355-4367.	1.3	41
12	The DAISY gene from Arabidopsis encodes a fatty acid elongase condensing enzyme involved in the biosynthesis of aliphatic suberin in roots and the chalazal-micropyle region of seeds. Plant Journal, 2009, 57, 80-95.	2.8	177
13	The grapevine transcription factor WRKY2 influences the lignin pathway and xylem development in tobacco. Plant Molecular Biology, 2010, 72, 215-234.	2.0	141
14	A feruloyl transferase involved in the biosynthesis of suberin and suberin-associated wax is required for maturation and sealing properties of potato periderm. Plant Journal, 2010, 62, 277-290.	2.8	120
15	Unraveling ferulate role in suberin and periderm biology by reverse genetics. Plant Signaling and Behavior, 2010, 5, 953-958.	1.2	24
16	Suberized Cell Walls of Cork from Cork Oak Differ from Other Species. Microscopy and Microanalysis, 2010, 16, 569-575.	0.2	26
17	Transport barriers made of cutin, suberin and associated waxes. Trends in Plant Science, 2010, 15, 546-553.	4.3	308
18	Phenylpropanoid Biosynthesis. Molecular Plant, 2010, 3, 2-20.	3.9	2,042

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19	Acyl-Lipid Metabolism. The Arabidopsis Book, 2010, 8, e0133.	0.5	287
20	Silicon enhances suberization and lignification in roots of rice ( <i>Oryza sativa</i> ). Journal of Experimental Botany, 2011, 62, 2001-2011.	2.4	210
23	Quercus. , 2011, , 89-129.		46
24	Temperature stress effects in <i>Quercus suber</i> leaf metabolism. Journal of Plant Physiology, 2011, 168, 1729-1734.	1.6	26
25	Suberin research in the genomics era—New interest for an old polymer. Plant Science, 2011, 180, 399-413.	1.7	185
26	Annexins. New Phytologist, 2011, 189, 40-53.	3.5	173
27	A potato skin SSH library yields new candidate genes for suberin biosynthesis and periderm formation. Planta, 2011, 233, 933-945.	1.6	39
28	Identification of GPAT acyltransferases in cork oak. BMC Proceedings, 2011, 5, .	1.8	4
29	Proteins associated with cork formation in <i>Quercus suber</i> L. stem tissues. Journal of Proteomics, 2011, 74, 1266-1278.	1.2	35
30	Metabolic engineering of novel lignin in biomass crops. New Phytologist, 2012, 196, 978-1000.	3.5	338
32	Reference Gene Selection for Quantitative Real-time PCR Normalization in <i>Quercus suber</i> . PLoS ONE, 2012, 7, e35113.	1.1	102
33	Genomics of Fagaceae. Tree Genetics and Genomes, 2012, 8, 583-610.	0.6	109
34	Arabidopsis ACA7, encoding a putative auto-regulated Ca <sup>2+</sup> -ATPase, is required for normal pollen development. Plant Cell Reports, 2012, 31, 651-659.	2.8	43
35	Molecular characterization of <i>Quercus suber</i> MYB1, a transcription factor up-regulated in cork tissues. Journal of Plant Physiology, 2013, 170, 172-178.	1.6	31
37	Acyl-Lipid Metabolism. The Arabidopsis Book, 2013, 11, e0161.	0.5	974
38	Identification of nitrogen responsive genes in poplar roots grown under two contrasting nitrogen levels. Plant Root, 2014, 8, 42-54.	0.3	2
39	miRNA profiling in leaf and cork tissues of <i>Quercus suber</i> reveals novel miRNAs and tissue-specific expression patterns. Tree Genetics and Genomes, 2014, 10, 721-737.	0.6	20
40	At<sc>MYB</sc>41 activates ectopic suberin synthesis and assembly in multiple plant species and cell types. Plant Journal, 2014, 80, 216-229.	2.8	172

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41	A comprehensive assessment of the transcriptome of cork oak ( <i>Quercus suber</i> ) through EST sequencing. <i>BMC Genomics</i> , 2014, 15, 371.	1.2	53
42	Comparison of good- and bad-quality cork: application of high-throughput sequencing of phellogenic tissue. <i>Journal of Experimental Botany</i> , 2014, 65, 4887-4905.	2.4	42
43	Microarray analysis of laser-microdissected tissues indicates the biosynthesis of suberin in the outer part of roots during formation of a barrier to radial oxygen loss in rice ( <i>Oryza sativa</i> ). <i>Journal of Experimental Botany</i> , 2014, 65, 4795-4806.	2.4	83
44	Relationship Between Tuber Storage Proteins and Tuber Powdery Scab Resistance in Potato. <i>American Journal of Potato Research</i> , 2014, 91, 233-245.	0.5	9
45	Physiological and transcriptional analyses of developmental stages along sugarcane leaf. <i>BMC Plant Biology</i> , 2015, 15, 300.	1.6	64
46	How tree roots respond to drought. <i>Frontiers in Plant Science</i> , 2015, 6, 547.	1.7	520
47	Examination of the Abscission-Associated Transcriptomes for Soybean, Tomato, and Arabidopsis Highlights the Conserved Biosynthesis of an Extensible Extracellular Matrix and Boundary Layer. <i>Frontiers in Plant Science</i> , 2015, 6, 1109.	1.7	38
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49	Suberin: biosynthesis, regulation, and polymer assembly of a protective extracellular barrier. <i>Plant Cell Reports</i> , 2015, 34, 573-586.	2.8	225
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56	Proteome Analyses of Soil Bacteria Grown in the Presence of Potato Suberin, a Recalcitrant Biopolymer. <i>Microbes and Environments</i> , 2016, 31, 418-426.	0.7	6
57	Asymmetrical development of root endodermis and exodermis in reaction to abiotic stresses. <i>Annals of Botany</i> , 2016, 118, 667-674.	1.4	54
58	MYB107 and MYB9 Homologs Regulate Suberin Deposition in Angiosperms. <i>Plant Cell</i> , 2016, 28, 2097-2116.	3.1	151
59	Silencing of the potato <i>StNAC103</i> gene enhances the accumulation of suberin polyester and associated wax in tuber skin. <i>Journal of Experimental Botany</i> , 2016, 67, 5415-5427.	2.4	56
60	Toughing It Out – Disease-Resistant Potato Mutants Have Enhanced Tuber Skin Defenses. <i>Phytopathology</i> , 2016, 106, 474-483.	1.1	41

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62	Characterization of the watercress ( <i>Nasturtium officinale</i> R. Br.; Brassicaceae) transcriptome using RNASeq and identification of candidate genes for important phytonutrient traits linked to human health. <i>BMC Genomics</i> , 2016, 17, 378.	1.2	33
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67	The Role of Hybridization on the Adaptive Potential of Mediterranean Sclerophyllous Oaks: The Case of the <i>Quercus ilex</i> x <i>Q. suber</i> Complex. <i>Tree Physiology</i> , 2017, , 239-260.	0.9	7
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70	A molecular framework to study periderm formation in <i>Arabidopsis</i> . <i>New Phytologist</i> , 2018, 219, 216-229.	3.5	78
71	Suberized transport barriers in <i>Arabidopsis</i> , barley and rice roots: From the model plant to crop species. <i>Journal of Plant Physiology</i> , 2018, 227, 75-83.	1.6	79
72	Cell Wall Reinforcement in the Potato Tuber Periderm After Crop Treatment with Potassium Phosphite. <i>Potato Research</i> , 2018, 61, 19-29.	1.2	11
73	Reconstructing the suberin pathway in poplar by chemical and transcriptomic analysis of bark tissues. <i>Tree Physiology</i> , 2018, 38, 340-361.	1.4	51
74	A comparative transcriptomic approach to understanding the formation of cork. <i>Plant Molecular Biology</i> , 2018, 96, 103-118.	2.0	35
75	Abscisic acid induces differential expression of genes involved in wound-induced suberization in postharvest tomato fruit. <i>Journal of Integrative Agriculture</i> , 2018, 17, 2670-2682.	1.7	16
76	Transcriptional profiling of cork oak phellogenic cells isolated by laser microdissection. <i>Planta</i> , 2018, 247, 317-338.	1.6	46
77	ChIP-Seq reveals that QsMYB1 directly targets genes involved in lignin and suberin biosynthesis pathways in cork oak ( <i>Quercus suber</i> ). <i>BMC Plant Biology</i> , 2018, 18, 198.	1.6	50
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80	Different clonal responses to cypress canker disease based on transcription of suberin-related genes and bark carbohydratesâ€™ content. Trees - Structure and Function, 2018, 32, 1707-1722.	0.9	7
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98	Transcriptomic analysis of cork during seasonal growth highlights regulatory and developmental processes from phellogen to phellem formation. <i>Scientific Reports</i> , 2021, 11, 12053.	1.6	13
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103	Differential DNA Methylation Patterns Are Related to Phellogen Origin and Quality of <i>Quercus suber</i> Cork. <i>PLoS ONE</i> , 2017, 12, e0169018.	1.1	31
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106	microRNA-Mediated Regulation of Plant Vascular Development and Secondary Growth. <i>Concepts and Strategies in Plant Sciences</i> , 2020, , 143-168.	0.6	1
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108	Characterization of a pericarp browning related LACCASE 14-4 from longan fruit with a focus on (epi)catechin oxidative polymerization. <i>Postharvest Biology and Technology</i> , 2022, 185, 111802.	2.9	8
109	Revealing the Genetic Components Responsible for the Unique Photosynthetic Stem Capability of the Wild Almond <i>Prunus arabica</i> (Olivier) Meikle. <i>Frontiers in Plant Science</i> , 2021, 12, 779970.	1.7	5
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111	The Making of Plant Armor: The Periderm. <i>Annual Review of Plant Biology</i> , 2022, 73, 405-432.	8.6	30
112	Genome-Wide Characterization and Abiotic Stresses Expression Analysis of Annexin Family Genes in Poplar. <i>International Journal of Molecular Sciences</i> , 2022, 23, 515.	1.8	3
113	Translational profile of developing phellem cells in <i>Arabidopsis thaliana</i> roots. <i>Plant Journal</i> , 2022, 110, 899-915.	2.8	9
114	Suberin Biosynthesis, Assembly, and Regulation. <i>Plants</i> , 2022, 11, 555.	1.6	44
115	Spatiotemporal development of suberized barriers in cork oak taproots. <i>Tree Physiology</i> , 2022, 42, 1269-1285.	1.4	4
129	Synthesis of C <sub>20</sub> –C <sub>38</sub> Fatty Acids in Plant Tissues. <i>International Journal of Molecular Sciences</i> , 2022, 23, 4731.	1.8	6

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130	The metabolic and proteomic repertoires of periderm tissue in skin of the reticulated Sikkim cucumber fruit. <i>Horticulture Research</i> , 2022, 9, .	2.9	10
131	<i>Quercus suber</i> Transcriptome Analyses: Identification of Genes and SNPs Related to Cork Quality. , 2022, 11, .		2
133	MYB1R1 and MYC2 Regulate Î³-3 Fatty Acid Desaturase Involved in ABA-Mediated Suberization in the Russet Skin of a Mutant of "Dangshansuli"™ ( <i>Pyrus bretschneideri</i> Rehd.). <i>Frontiers in Plant Science</i> , 0, 13, .	1.7	6
134	Effect of climate on cork-ring width and density of <i>Quercus suber</i> L. in Southern Portugal. <i>Trees - Structure and Function</i> , 2022, 36, 1711-1720.	0.9	6
135	Knockout of <i>MITOGEN-ACTIVATED PROTEIN KINASE 3</i> causes barley root resistance against <i>Fusarium graminearum</i> . <i>Plant Physiology</i> , 2022, 190, 2847-2867.	2.3	7
136	<i>Populus</i> — <i>canescens</i> root suberization in reaction to osmotic and salt stress is limited to the developing younger root tip region. <i>Physiologia Plantarum</i> , 2022, 174, .	2.6	6
137	Cork Development: What Lies Within. <i>Plants</i> , 2022, 11, 2671.	1.6	7
138	Beyond width and density: stable carbon and oxygen isotopes in cork-rings provide insights of physiological responses to water stress in <i>Quercus suber</i> L. <i>PeerJ</i> , 0, 10, e14270.	0.9	1
139	Foxtail millet MYB-like transcription factor SiMYB16 confers salt tolerance in transgenic rice by regulating phenylpropane pathway. <i>Plant Physiology and Biochemistry</i> , 2023, 195, 310-321.	2.8	9
140	The application of fatty acid synthesis inhibitors regulates the suberin biosynthesis in the exocarp of the russet mutant of "Dangshansuli"™ ( <i>Pyrus bretschneideri</i> Rehd.). <i>Acta Physiologiae Plantarum</i> , 2023, 45, .	1.0	0
141	The D165H Polymorphism of QiMYB-like-1 Is Linked to Interactions between Tannin Accumulation, Herbivory and Biogeographical Determinants of <i>Quercus ilex</i> . <i>International Journal of Molecular Sciences</i> , 2023, 24, 151.	1.8	0
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