JÃ¶rg Bohlmann

List of Publications by Year
in descending order

227	23,371	85	143
papers	citations	h-index	g-index
		4658	9345

232	232	232	16972
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	The Norway spruce genome sequence and conifer genome evolution. Nature, 2013, 497, 579-584.	27.8	1,303
2	The family of terpene synthases in plants: a midâ€size family of genes for specialized metabolism that is highly diversified throughout the kingdom. Plant Journal, 2011, 66, 212-229.	5.7	1,068
3	Genes, enzymes and chemicals of terpenoid diversity in the constitutive and induced defence of conifers against insects and pathogens*. New Phytologist, 2006, 170, 657-675.	7.3	593
4	Methyl Jasmonate Induces Traumatic Resin Ducts, Terpenoid Resin Biosynthesis, and Terpenoid Accumulation in Developing Xylem of Norway Spruce Stems. Plant Physiology, 2002, 129, 1003-1018.	4.8	462
5	Terpenoid biomaterials. Plant Journal, 2008, 54, 656-669.	5.7	423
6	(E)-Î²-Ocimene and Myrcene Synthase Genes of Floral Scent Biosynthesis in Snapdragon: Function and Expression of Three Terpene Synthase Genes of a New Terpene Synthase Subfamily. Plant Cell, 2003, 15, 1227-1241.	6.6	397
7	Functional Annotation, Genome Organization and Phylogeny of the Grapevine (Vitis vinifera) Terpene Synthase Gene Family Based on Genome Assembly, FLcDNA Cloning, and Enzyme Assays. BMC Plant Biology, 2010, 10, 226.	3.6	390
8	Induction of Volatile Terpene Biosynthesis and Diurnal Emission by Methyl Jasmonate in Foliage of Norway Spruce. Plant Physiology, 2003, 132, 1586-1599.	4.8	381
9	Functional Characterization of Nine Norway Spruce TPS Genes and Evolution of Gymnosperm Terpene Synthases of the TPS-d Subfamily. Plant Physiology, 2004, 135, 1908-1927.	4.8	369
10	Sesquiterpene Synthases from Grand Fir (Abies grandis). Journal of Biological Chemistry, 1998, 273, 2078-2089.	3.4	362
11	Assembling the 20 Gb white spruce (<i>Picea glauca</i>) genome from whole-genome shotgun sequencing data. Bioinformatics, 2013, 29, 1492-1497.	4.1	356
12	Diterpene resin acids in conifers. Phytochemistry, 2006, 67, 2415-2423.	2.9	284
13	Insect-Induced Conifer Defense. White Pine Weevil and Methyl Jasmonate Induce Traumatic Resinosis, de Novo Formed Volatile Emissions, and Accumulation of Terpenoid Synthase and Putative Octadecanoid Pathway Transcripts in Sitka Spruce. Plant Physiology, 2005, 137, 369-382.	4.8	262
14	Draft genome of the mountain pine beetle, Dendroctonus ponderosae Hopkins, a major forest pest. Genome Biology, 2013, 14, R27.	9.6	260
15	Global transcript profiling of primary stems from Arabidopsis thaliana identifies candidate genes for missing links in lignin biosynthesis and transcriptional regulators of fiber differentiation. Plant Journal, 2005, 42, 618-640.	5.7	254
16	Terpenoid Biosynthesis and Specialized Vascular Cells of Conifer Defense. Journal of Integrative Plant Biology, 2010, 52, 86-97.	8.5	254
17	Efficacy of tree defense physiology varies with bark beetle population density: a basis for positive feedback in eruptive species. Canadian Journal of Forest Research, 2011, 41, 1174-1188.	1.7	250
18	Pine monoterpenes and pine bark beetles: a marriage of convenience for defense and chemical communication. Phytochemistry Reviews, 2006, 5, 143-178.	6.5	233

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19	Transgenic, non-isoprene emitting poplars donâ€™t like it hot. Plant Journal, 2007, 51, 485-499.	5.7	229
20	Bacteria Associated with a Tree-Killing Insect Reduce Concentrations of Plant Defense Compounds. Journal of Chemical Ecology, 2013, 39, 1003-1006.	1.8	227
21	Conifer defence against insects: microarray gene expression profiling of Sitka spruce (Picea) Tj ETQq1 1 0.784314 rgBT /Overlock 10 transcriptome. Plant, Cell and Environment, 2006, 29, 1545-1570.	5.7	221
22	Forest tent caterpillars (Malacosoma disstria) induce local and systemic diurnal emissions of terpenoid volatiles in hybrid poplar (Populus trichocarpaâ€ƒÃ—â€ƒdeltoides): cDNA cloning, functional characterization, and patterns of gene expression of (â˜’)-germacr. Plant Journal, 2004, 37, 603-616.	5.7	220
23	Genome and transcriptome analyses of the mountain pine beetle-fungal symbiont <i>Grosmannia clavigera</i> , a lodgepole pine pathogen. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 2504-2509.	7.1	218
24	Antennal transcriptome analysis of the chemosensory gene families in the tree killing bark beetles, Ips typographus and Dendroctonus ponderosae (Coleoptera: Curculionidae: Scolytinae). BMC Genomics, 2013, 14, 198.	2.8	216
25	Monoterpene Synthases from Grand Fir (Abies grandis). Journal of Biological Chemistry, 1997, 272, 21784-21792.	3.4	210
26	Characterization of a Root-Specific Arabidopsis Terpene Synthase Responsible for the Formation of the Volatile Monoterpene 1,8-Cineole. Plant Physiology, 2004, 135, 1956-1966.	4.8	207
27	Vitis vinifera terpenoid cyclases: functional identification of two sesquiterpene synthase cDNAs encoding (+)-valencene synthase and (â˜’)-germacrene D synthase and expression of mono- and sesquiterpene synthases in grapevine flowers and berries. Phytochemistry, 2004, 65, 2649-2659.	2.9	205
28	The Transcriptional Response of Hybrid Poplar (Populus trichocarpa x P. deltoids) to Infection by Melampsora medusae Leaf Rust Involves Induction of Flavonoid Pathway Genes Leading to the Accumulation of Proanthocyanidins. Molecular Plant-Microbe Interactions, 2007, 20, 816-831.	2.6	205
29	Improved white spruce (<i>Picea glauca</i>) genome assemblies and annotation of large gene families of conifer terpenoid and phenolic defense metabolism. Plant Journal, 2015, 83, 189-212.	5.7	200
30	Transcriptome analysis based on next-generation sequencing of non-model plants producing specialized metabolites of biotechnological interest. Journal of Biotechnology, 2013, 166, 122-134.	3.8	196
31	Genomics of hybrid poplar (Populus trichocarpaÃ— deltoides) interacting with forest tent caterpillars (Malacosoma disstria): normalized and full-length cDNA libraries, expressed sequence tags, and a cDNA microarray for the study of insect-induced defences. Molecular Ecology, 2006, 15, 1275-1297.	3.9	183
32	Terpene synthases from Cannabis sativa. PLoS ONE, 2017, 12, e0173911.	2.5	183
33	Robust simple sequence repeat markers for spruce (Picea spp.) from expressed sequence tags. Theoretical and Applied Genetics, 2004, 109, 1283-1294.	3.6	181
34	Functional plasticity of paralogous diterpene synthases involved in conifer defense. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 1085-1090.	7.1	178
35	Loblolly pine abietadienol/abietadienal oxidase PtAO (CYP720B1) is a multifunctional, multisubstrate cytochrome P450 monooxygenase. Proceedings of the National Academy of Sciences of the United States of America, 2005, 102, 8060-8065.	7.1	172
36	Traumatic resin defense in Norway spruce (Picea abies): methyl jasmonate-induced terpene synthase gene expression, and cDNA cloning and functional characterization of (+)-3-carene synthase. Plant Molecular Biology, 2003, 51, 119-133.	3.9	171

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37	Proteome analysis of early somatic embryogenesis inPicea glauca. Proteomics, 2005, 5, 461-473.	2.2	166
38	Slow but not low: genomic comparisons reveal slower evolutionary rate and higher dN/dS in conifers compared to angiosperms. BMC Evolutionary Biology, 2012, 12, 8.	3.2	164
39	Insights into Conifer Giga-Genomes. Plant Physiology, 2014, 166, 1724-1732.	4.8	164
40	Dirigent Proteins in Conifer Defense: Gene Discovery, Phylogeny, and Differential Wound- and Insect-induced Expression of a Family of DIR and DIR-like Genes in Spruce (Picea spp.). Plant Molecular Biology, 2006, 60, 21-40.	3.9	160
41	The Molecular Basis for Wine Grape Quality-A Volatile Subject. Science, 2006, 311, 804-805.	12.6	158
42	Cannabis glandular trichomes alter morphology and metabolite content during flower maturation. Plant Journal, 2020, 101, 37-56.	5.7	158
43	Terpenes in Cannabis sativa â€“ From plant genome to humans. Plant Science, 2019, 284, 67-72.	3.6	157
44	Gene Discovery of Modular Diterpene Metabolism in Nonmodel Systems Â Â. Plant Physiology, 2013, 162, 1073-1091.	4.8	154
45	A specialized <scp>ABC</scp> efflux transporter <scp>G</scp>c<scp>ABC</scp>â€<scp>G</scp>1 confers monoterpene resistance to <i><scp>G</scp>rosmannia clavigera</i>, a bark beetleâ€associated fungal pathogen of pine trees. New Phytologist, 2013, 197, 886-898.	7.3	152
46	Discovery and functional characterization of two diterpene synthases for sclareol biosynthesis in Salvia sclarea(L.) and their relevance for perfume manufacture. BMC Plant Biology, 2012, 12, 119.	3.6	151
47	A Common Fungal Associate of the Spruce Bark Beetle Metabolizes the Stilbene Defenses of Norway Spruce Â Â Â. Plant Physiology, 2013, 162, 1324-1336.	4.8	150
48	Evolution of Diterpene Metabolism: Sitka Spruce CYP720B4 Catalyzes Multiple Oxidations in Resin Acid Biosynthesis of Conifer Defense against Insects Â Â Â. Plant Physiology, 2011, 157, 1677-1695.	4.8	149
49	Isolation of high-quality RNA from gymnosperm and angiosperm trees. BioTechniques, 2004, 36, 821-824.	1.8	148
50	Insect Attack and Wounding Induce Traumatic Resin Duct Development and Gene Expression of (â€”)-Pinene Synthase in Sitka Spruce. Plant Physiology, 2003, 133, 368-378.	4.8	144
51	Mono and diterpene production inEscherichia coli. Biotechnology and Bioengineering, 2004, 87, 200-212.	3.3	141
52	Use of Ecotilling as an efficient SNP discovery tool to survey genetic variation in wild populations of Populus trichocarpa. Molecular Ecology, 2006, 15, 1367-1378.	3.9	140
53	Herbivore-Induced Defense Response in a Model Legume. Two-Spotted Spider Mites Induce Emission of (<i>E</i>)-<i>Î²</i>-Ocimene and Transcript Accumulation of (<i>E</i>)-<i>Î²</i>-Ocimene Synthase in <i>Lotus japonicus</i> Â. Plant Physiology, 2004, 135, 1976-1983.	4.8	139
54	Oleoresin defenses in conifers: chemical diversity, terpene synthases and limitations of oleoresin defense under climate change. New Phytologist, 2019, 224, 1444-1463.	7.3	139

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55	Terpenoid Secondary Metabolism in Arabidopsis thaliana: cDNA Cloning, Characterization, and Functional Expression of a Myrcene/(E)-Î²-Ocimene Synthase. Archives of Biochemistry and Biophysics, 2000, 375, 261-269.	3.0	137
56	Manoyl Oxide (13R), the Biosynthetic Precursor of Forskolin, Is Synthesized in Specialized Root Cork Cells in <i>Coleus forskohlii</i>. Plant Physiology, 2014, 164, 1222-1236.	4.8	135
57	Functional identification of AtTPS03 as (E)-Î²-ocimene synthase: a monoterpene synthase catalyzing jasmonate- and wound-induced volatile formation in Arabidopsis thaliana. Planta, 2003, 216, 745-751.	3.2	134
58	Expanding the Landscape of Diterpene Structural Diversity through Stereochemically Controlled Combinatorial Biosynthesis. Angewandte Chemie - International Edition, 2016, 55, 2142-2146.	13.8	134
59	Plant diterpene synthases: exploring modularity and metabolic diversity for bioengineering. Trends in Biotechnology, 2015, 33, 419-428.	9.3	133
60	cDNA Cloning, Characterization, and Functional Expression of Four New Monoterpene Synthase Members of the Tpsd Gene Family from Grand Fir (Abies grandis). Archives of Biochemistry and Biophysics, 1999, 368, 232-243.	3.0	130
61	De novo genome sequence assembly of a filamentous fungus using Sanger, 454 and Illumina sequence data. Genome Biology, 2009, 10, R94.	9.6	130
62	Synthetic biosystems for the production of high-value plant metabolites. Trends in Biotechnology, 2012, 30, 127-131.	9.3	128
63	Sandalwood Fragrance Biosynthesis Involves Sesquiterpene Synthases of Both the Terpene Synthase (TPS)-a and TPS-b Subfamilies, including Santalene Synthases. Journal of Biological Chemistry, 2011, 286, 17445-17454.	3.4	127
64	Functional identification and differential expression of 1-deoxy-d-xylulose 5-phosphate synthase in induced terpenoid resin formation of Norway spruce (Picea abies). Plant Molecular Biology, 2007, 65, 243-257.	3.9	126
65	Identification of Vitis vinifera (âˆ’)-Î±-terpineol synthase by in silico screening of full-length cDNA ESTs and functional characterization of recombinant terpene synthase. Phytochemistry, 2004, 65, 1223-1229.	2.9	122
66	BIOSYNTHESIS OF CONIFEROPHAGOUS BARK BEETLE PHEROMONES AND CONIFER ISOPRENOIDS: EVOLUTIONARY PERSPECTIVE AND SYNTHESIS. Canadian Entomologist, 2000, 132, 697-753.	0.8	120
67	Transcriptome mining, functional characterization, and phylogeny of a large terpene synthase gene family in spruce (Piceaspp.). BMC Plant Biology, 2011, 11, 43.	3.6	120
68	Evolution of Conifer Diterpene Synthases: Diterpene Resin Acid Biosynthesis in Lodgepole Pine and Jack Pine Involves Monofunctional and Bifunctional Diterpene Synthases Â Â. Plant Physiology, 2013, 161, 600-616.	4.8	118
69	Biosynthesis of Sandalwood Oil: Santalum album CYP76F Cytochromes P450 Produce Santalols and Bergamotol. PLoS ONE, 2013, 8, e75053.	2.5	117
70	Dirigent proteins in conifer defense II: Extended gene discovery, phylogeny, and constitutive and stress-induced gene expression in spruce (Picea spp.)â˜†. Phytochemistry, 2007, 68, 1975-1991.	2.9	116
71	An integrated genomic, proteomic and biochemical analysis of (+)â€3â€carene biosynthesis in Sitka spruce (<i>Picea sitchensis</i>) genotypes that are resistant or susceptible to white pine weevil. Plant Journal, 2011, 65, 936-948.	5.7	116
72	Population sequencing reveals clonal diversity and ancestral inbreeding in the grapevine cultivar Chardonnay. PLoS Genetics, 2018, 14, e1007807.	3.5	116

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73	Monoterpene-induced molecular responses in Arabidopsis thaliana. Phytochemistry, 2008, 69, 1838-1849.	2.9	113
74	A conifer genomics resource of 200,000 spruce (Picea spp.) ESTs and 6,464 high-quality, sequence-finished full-length cDNAs for Sitka spruce (Picea sitchensis). BMC Genomics, 2008, 9, 484.	2.8	113
75	Biosynthesis of the Major Tetrahydroxystilbenes in Spruce, Astringin and Isorhapontin, Proceeds via Resveratrol and Is Enhanced by Fungal Infection Â Â. Plant Physiology, 2011, 157, 876-890.	4.8	112
76	Biosynthesis of wine aroma: transcript profiles of hydroxymethylbutenyl diphosphate reductase, geranyl diphosphate synthase, and linalool/nerolidol synthase parallel monoterpenol glycoside accumulation in GewÃ¼rztraminer grapes. Planta, 2012, 236, 919-929.	3.2	112
77	Global monitoring of autumn gene expression within and among phenotypically divergent populations of Sitka spruce (<i>Picea sitchensis</i>). New Phytologist, 2008, 178, 103-122.	7.3	111
78	Wound-Induced Terpene Synthase Gene Expression in Sitka Spruce That Exhibit Resistance or Susceptibility to Attack by the White Pine Weevil Â. Plant Physiology, 2006, 140, 1009-1021.	4.8	106
79	Targeted proteomics using selected reaction monitoring reveals the induction of specific terpene synthases in a multiâ€level study of methyl jasmonateâ€treated Norway spruce (<i>Picea abies</i>). Plant Journal, 2009, 60, 1015-1030.	5.7	106
80	Impact of <i>Salmonella</i> Infection on Host Hormone Metabolism Revealed by Metabolomics. Infection and Immunity, 2011, 79, 1759-1769.	2.2	104
81	Contribution of isopentenyl phosphate to plant terpenoid metabolism. Nature Plants, 2018, 4, 721-729.	9.3	100
82	Identification and Functional Characterization of Monofunctional<i>ent</i>-Copalyl Diphosphate and<i>ent</i>-Kaurene Synthases in White Spruce Reveal Different Patterns for Diterpene Synthase Evolution for Primary and Secondary Metabolism in Gymnosperms. Plant Physiology, 2010, 152, 1197-1208.	4.8	99
83	The bouquet of grapevine (<i>Vitis vinifera</i> L. cv. Cabernet Sauvignon) flowers arises from the biosynthesis of sesquiterpene volatiles in pollen grains. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 7245-7250.	7.1	97
84	Tigmint: correcting assembly errors using linked reads from large molecules. BMC Bioinformatics, 2018, 19, 393.	2.6	97
85	Discovery, Biosynthesis and Stress-Related Accumulation of Dolabradiene-Derived Defenses in Maize. Plant Physiology, 2018, 176, 2677-2690.	4.8	94
86	Transcriptome and full-length cDNA resources for the mountain pine beetle, Dendroctonus ponderosae Hopkins, a major insect pest of pine forests. Insect Biochemistry and Molecular Biology, 2012, 42, 525-536.	2.7	93
87	Comparative transcriptome analysis of Arabidopsis thaliana infested by diamond back moth (Plutella) Tj ETQq1 1 0.784314 rgBT /Overlo Genomics, 2008, 9, 154.	2.8	90
88	SNP discovery, gene diversity, and linkage disequilibrium in wild populations of Populus tremuloides. Tree Genetics and Genomes, 2012, 8, 821-829.	1.6	86
89	Toxicity of Pine Monoterpenes to Mountain Pine Beetle. Scientific Reports, 2017, 7, 8858.	3.3	85
90	Laser microdissection of conifer stem tissues: Isolation and analysis of high quality RNA, terpene synthase enzyme activity and terpenoid metabolites from resin ducts and cambial zone tissue of white spruce (Picea glauca). BMC Plant Biology, 2010, 10, 106.	3.6	83

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91	Conifer defense against insects: Proteome analysis of Sitka spruce (Picea sitchensis) bark induced by mechanical wounding or feeding by white pine weevils (Pissodes strobi). Proteomics, 2007, 7, 248-270.	2.2	82
92	RNA-seq discovery, functional characterization, and comparison of sesquiterpene synthases from Solanum lycopersicum and Solanum habrochaites trichomes. Plant Molecular Biology, 2011, 77, 323-336.	3.9	80
93	Heartwoodâ€specific transcriptome and metabolite signatures of tropical sandalwood (<i>Santalum) Tj ETQq1 1 0.784314 rgBT /Overlo 289-299.	5.7	79
94	Purification and cDNA cloning of anthranilate synthase from Ruta graveolens: modes of expression and properties of native and recombinant enzymes. Plant Journal, 1995, 7, 491-501.	5.7	78
95	Diterpene resin acid biosynthesis in loblolly pine (Pinus taeda): Functional characterization of abietadiene/levopimaradiene synthase (PtTPS-LAS) cDNA and subcellular targeting of PtTPS-LAS and abietadienol/abietadienal oxidase (PtAO, CYP720B1). Phytochemistry, 2006, 67, 1572-1578.	2.9	78
96	Changes in anatomy and terpene chemistry in roots of Douglas-fir seedlings following treatment with methyl jasmonate. Tree Physiology, 2005, 25, 1075-1083.	3.1	77
97	Bifunctional cis-Abienol Synthase from Abies balsamea Discovered by Transcriptome Sequencing and Its Implications for Diterpenoid Fragrance Production. Journal of Biological Chemistry, 2012, 287, 12121-12131.	3.4	75
98	Ethylene in induced conifer defense: cDNA cloning, protein expression, and cellular and subcellular localization of 1-aminocyclopropane-1-carboxylate oxidase in resin duct and phenolic parenchyma cells. Planta, 2006, 224, 865-877.	3.2	74
99	Gene Discovery for Enzymes Involved in Limonene Modification or Utilization by the Mountain Pine Beetle-Associated Pathogen Grosmannia clavigera. Applied and Environmental Microbiology, 2014, 80, 4566-4576.	3.1	74
100	GENOMIC HARDWIRING AND PHENOTYPIC PLASTICITY OF TERPENOID-BASED DEFENSES IN CONIFERS. Journal of Chemical Ecology, 2004, 30, 2399-2418.	1.8	73
101	The versatility of the fungal cytochrome P450 monooxygenase system is instrumental in xenobiotic detoxification. Molecular Microbiology, 2011, 81, 1374-1389.	2.5	73
102	Responses of Bark Beetle-Associated Bacteria to Host Monoterpenes and Their Relationship to Insect Life Histories. Journal of Chemical Ecology, 2011, 37, 808-817.	1.8	73
103	Genomeâ€wide analysis of a land plantâ€specific <i>acyl:coenzymeA synthetase</i> (<i>ACS</i>) gene family in <i>Arabidopsis</i>, poplar, rice and <i>Physcomitrella</i>. New Phytologist, 2008, 179, 987-1003.	7.3	72
104	Isolation of cDNAs and functional characterisation of two multi-product terpene synthase enzymes from sandalwood, Santalum album L.. Archives of Biochemistry and Biophysics, 2008, 477, 121-130.	3.0	72
105	The genome and transcriptome of the pine saprophyte Ophiostoma piceae, and a comparison with the bark beetle-associated pine pathogen Grosmannia clavigera. BMC Genomics, 2013, 14, 373.	2.8	72
106	Flavan-3-ols in Norway Spruce: Biosynthesis, Accumulation, and Function in Response to Attack by the Bark Beetle-Associated Fungus <i>Ceratocystis polonica</i> Â Â Â Â. Plant Physiology, 2014, 164, 2107-2122.	4.8	72
107	Characterization of four terpene synthase cDNAs from methyl jasmonate-induced Douglas-fir, Pseudotsuga menziesii. Phytochemistry, 2005, 66, 1427-1439.	2.9	70
108	A physical map of the highly heterozygous Populus genome: integration with the genome sequence and genetic map and analysis of haplotype variation. Plant Journal, 2007, 50, 1063-1078.	5.7	70

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109	How the Mountain Pine Beetle (Dendroctonus ponderosae) Breached the Canadian Rocky Mountains. Molecular Biology and Evolution, 2014, 31, 1803-1815.	8.9	70
110	Analysis of 4,664 high-quality sequence-finished poplar full-length cDNA clones and their utility for the discovery of genes responding to insect feeding. BMC Genomics, 2008, 9, 57.	2.8	68
111	ntEdit: scalable genome sequence polishing. Bioinformatics, 2019, 35, 4430-4432.	4.1	67
112	Poplar defense against insect herbivoresThis review is one of a selection of papers published in the Special Issue on Poplar Research in Canada.. Canadian Journal of Botany, 2007, 85, 1111-1126.	1.1	65
113	Diterpene synthases of the biosynthetic system of medicinally active diterpenoids in <i>Marrubium vulgare</i>. Plant Journal, 2014, 79, 914-927.	5.7	62
114	Global and comparative proteomic profiling of overwintering and developing mountain pine beetle, Dendroctonus ponderosae (Coleoptera: Curculionidae), larvae. Insect Biochemistry and Molecular Biology, 2012, 42, 890-901.	2.7	61
115	CYP345E2, an antenna-specific cytochrome P450 from the mountain pine beetle, Dendroctonus ponderosae Hopkins, catalyses the oxidation of pine host monoterpene volatiles. Insect Biochemistry and Molecular Biology, 2013, 43, 1142-1151.	2.7	61
116	Transcriptional responses of Arabidopsis thaliana to chewing and sucking insect herbivores. Frontiers in Plant Science, 2014, 5, 565.	3.6	61
117	Microarray expression profiling and functional characterization of AtTPS genes: Duplicated Arabidopsis thaliana sesquiterpene synthase genes At4g13280 and At4g13300 encode root-specific and wound-inducible (Z)-Î³-bisabolene synthases. Archives of Biochemistry and Biophysics, 2006, 448, 104-116.	3.0	60
118	Multiple genes recruited from hormone pathways partition maize diterpenoid defences. Nature Plants, 2019, 5, 1043-1056.	9.3	60
119	Pine terpenoid defences in the mountain pine beetle epidemic and in other conifer pest interactions: specialized enemies are eating holes into a diverse, dynamic and durable defence system. Tree Physiology, 2012, 32, 943-945.	3.1	59
120	Terpenoid metabolite profiling in Sitka spruce identifies association of dehydroabietic acid, (+)-3-carene, and terpinolene with resistance against white pine weevil. Botany, 2010, 88, 810-820.	1.0	57
121	Gene genealogies reveal cryptic species and host preferences for the pine fungal pathogen<i>Grosmannia clavigera</i>. Molecular Ecology, 2011, 20, 2581-2602.	3.9	57
122	Transcriptome resources and functional characterization of monoterpene synthases for two host species of the mountain pine beetle, lodgepole pine (Pinus contorta) and jack pine (Pinus banksiana). BMC Plant Biology, 2013, 13, 80.	3.6	57
123	Disentangling Detoxification: Gene Expression Analysis of Feeding Mountain Pine Beetle Illuminates Molecular-Level Host Chemical Defense Detoxification Mechanisms. PLoS ONE, 2013, 8, e77777.	2.5	57
124	Targeted isolation, sequence assembly and characterization of two white spruce (Picea glauca) BAC clones for terpenoid synthase and cytochrome P450 genes involved in conifer defence reveal insights into a conifer genome. BMC Plant Biology, 2009, 9, 106.	3.6	55
125	Biosynthesis of the psychotropic plant diterpene salvinorin A: Discovery and characterization of the <i>Salvia divinorum</i> clerodienyl diphosphate synthase. Plant Journal, 2017, 89, 885-897.	5.7	55
126	Aminocyclopropane Carboxylic Acid Synthase Is a Regulated Step in Ethylene-Dependent Induced Conifer Defense. Full-Length cDNA Cloning of a Multigene Family, Differential Constitutive, and Wound- and Insect-Induced Expression, and Cellular and Subcellular Localization in Spruce and Douglas Fir. Plant Physiology, 2007, 143, 410-424.	4.8	54

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127	The Primary Diterpene Synthase Products of Picea abies Levopimaradiene/Abietadiene Synthase (PaLAS) Are Epimers of a Thermally Unstable Diterpenol. Journal of Biological Chemistry, 2011, 286, 21145-21153.	3.4	52
128	Expression of the Î²â€glucosidase gene <i>PgÎ²gluâ€1</i> underpins natural resistance of white spruce against spruce budworm. Plant Journal, 2015, 81, 68-80.	5.7	52
129	Genetic elucidation of interconnected antibiotic pathways mediating maize innate immunity. Nature Plants, 2020, 6, 1375-1388.	9.3	52
130	Terpene Synthases and Terpene Variation in <i>Cannabis sativa</i>. Plant Physiology, 2020, 184, 130-147.	4.8	52
131	Analysis of the Poplar Phloem Proteome and Its Response to Leaf Wounding. Journal of Proteome Research, 2009, 8, 2341-2350.	3.7	50
132	Poplar defense against insects: genome analysis, fullâ€length cDNA cloning, and transcriptome and protein analysis of the poplar Kunitzâ€type protease inhibitor family. New Phytologist, 2009, 184, 865-884.	7.3	49
133	Complete Mitochondrial Genome of a Gymnosperm, Sitka Spruce (Picea sitchensis), Indicates a Complex Physical Structure. Genome Biology and Evolution, 2020, 12, 1174-1179.	2.5	49
134	Transcriptome profiles of hybrid poplar (<i>Populus trichocarpa</i>â€ƒÃ—â€ƒ<i>deltoides</i>) reveal rapid changes in undamaged, systemic sink leaves after simulated feeding by forest tent caterpillar (<i>Malacosoma disstria</i>). New Phytologist, 2010, 188, 787-802.	7.3	48
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