

# Mark Lange

## List of Publications by Year in descending order

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80  
papers

9,075  
citations

117453

34  
h-index

66788

78  
g-index

96  
all docs

96  
docs citations

96  
times ranked

10481  
citing authors

#	ARTICLE	IF	CITATIONS
1	A Draft Sequence of the Rice Genome ( <i>Oryza sativa</i> L. ssp. japonica). <i>Science</i> , 2002, 296, 92-100.	6.0	2,866
2	Isoprenoid biosynthesis: The evolution of two ancient and distinct pathways across genomes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2000, 97, 13172-13177.	3.3	720
3	Potential of metabolomics as a functional genomics tool. <i>Trends in Plant Science</i> , 2004, 9, 418-425.	4.3	685
4	Proteomic survey of metabolic pathways in rice. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2002, 99, 11969-11974.	3.3	386
5	A family of transketolases that directs isoprenoid biosynthesis via a mevalonate-independent pathway. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1998, 95, 2100-2104.	3.3	351
6	Transcriptional regulators of stamen development in <i>Arabidopsis</i> identified by transcriptional profiling. <i>Plant Journal</i> , 2006, 46, 984-1008.	2.8	299
7	Probing essential oil biosynthesis and secretion by functional evaluation of expressed sequence tags from mint glandular trichomes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2000, 97, 2934-2939.	3.3	292
8	A proposed framework for the description of plant metabolomics experiments and their results. <i>Nature Biotechnology</i> , 2004, 22, 1601-1606.	9.4	283
9	Genome organization in <i>Arabidopsis thaliana</i> : a survey for genes involved in isoprenoid and chlorophyll metabolism. <i>Plant Molecular Biology</i> , 2003, 51, 925-948.	2.0	240
10	Metabolic engineering of plant monoterpenes, sesquiterpenes and diterpenes—current status and future opportunities. <i>Plant Biotechnology Journal</i> , 2013, 11, 169-196.	4.1	169
11	Isoprenoid Biosynthesis via a Mevalonate-Independent Pathway in Plants: Cloning and Heterologous Expression of 1-Deoxy-d-xylulose-5-phosphate Reductoisomerase from Peppermint. <i>Archives of Biochemistry and Biophysics</i> , 1999, 365, 170-174.	1.4	157
12	Terpenoid biosynthesis in trichomes—current status and future opportunities. <i>Plant Biotechnology Journal</i> , 2013, 11, 2-22.	4.1	146
13	The Evolution of Plant Secretory Structures and Emergence of Terpenoid Chemical Diversity. <i>Annual Review of Plant Biology</i> , 2015, 66, 139-159.	8.6	145
14	Improving peppermint essential oil yield and composition by metabolic engineering. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 16944-16949.	3.3	127
15	Minimum reporting standards for plant biology context information in metabolomic studies. <i>Metabolomics</i> , 2007, 3, 195-201.	1.4	116
16	A systems biology approach identifies the biochemical mechanisms regulating monoterpenoid essential oil composition in peppermint. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 2818-2823.	3.3	116
17	Open-Access Metabolomics Databases for Natural Product Research: Present Capabilities and Future Potential. <i>Frontiers in Bioengineering and Biotechnology</i> , 2015, 3, 22.	2.0	114
18	Isopentenyl diphosphate biosynthesis via a mevalonate-independent pathway: Isopentenyl monophosphate kinase catalyzes the terminal enzymatic step. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1999, 96, 13714-13719.	3.3	109

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19	Experimental and mathematical approaches to modeling plant metabolic networks. <i>Phytochemistry</i> , 2007, 68, 2351-2374.	1.4	101
20	PlantMetabolomics.org: A Web Portal for Plant Metabolomics Experiments. <i>Plant Physiology</i> , 2010, 152, 1807-1816.	2.3	93
21	Gene Networks Underlying Cannabinoid and Terpenoid Accumulation in Cannabis. <i>Plant Physiology</i> , 2019, 180, 1877-1897.	2.3	90
22	Assessing the Biosynthetic Capabilities of Secretory Glands in <i>Citrus</i> Peel. <i>Plant Physiology</i> , 2012, 159, 81-94.	2.3	82
23	Metabolomics as a Hypothesis-Generating Functional Genomics Tool for the Annotation of <i>Arabidopsis thaliana</i> Genes of "Unknown Function". <i>Frontiers in Plant Science</i> , 2012, 3, 15.	1.7	82
24	Draft Genome Sequence of <i>Mentha longifolia</i> and Development of Resources for Mint Cultivar Improvement. <i>Molecular Plant</i> , 2017, 10, 323-339.	3.9	79
25	Isoprenoid Biosynthesis. Metabolite Profiling of Peppermint Oil Gland Secretory Cells and Application to Herbicide Target Analysis. <i>Plant Physiology</i> , 2001, 127, 305-314.	2.3	76
26	Functional analysis of (4 <i>S</i> )-limonene synthase mutants reveals determinants of catalytic outcome in a model monoterpene synthase. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 3332-3337.	3.3	70
27	Comprehensive post-genomic data analysis approaches integrating biochemical pathway maps. <i>Phytochemistry</i> , 2005, 66, 413-451.	1.4	67
28	Integrative analysis of transcript and metabolite profiling data sets to evaluate the regulation of biochemical pathways during photomorphogenesis. <i>Archives of Biochemistry and Biophysics</i> , 2006, 448, 45-59.	1.4	66
29	Mathematical Modeling-Guided Evaluation of Biochemical, Developmental, Environmental, and Genotypic Determinants of Essential Oil Composition and Yield in Peppermint Leaves. <i>Plant Physiology</i> , 2010, 152, 2105-2119.	2.3	59
30	Patterns of Metabolite Changes Identified from Large-Scale Gene Perturbations in <i>Arabidopsis</i> Using a Genome-Scale Metabolic Network. <i>Plant Physiology</i> , 2015, 167, 1685-1698.	2.3	55
31	Integrative Approaches for the Identification and Localization of Specialized Metabolites in <i>Tripterygium</i> Roots. <i>Plant Physiology</i> , 2017, 173, 456-469.	2.3	47
32	Accurate mass-time tag library for LC/MS-based metabolite profiling of medicinal plants. <i>Phytochemistry</i> , 2013, 91, 187-197.	1.4	43
33	Abscisic acid-induced modulation of metabolic and redox control pathways in <i>Arabidopsis thaliana</i> . <i>Phytochemistry</i> , 2008, 69, 2899-2911.	1.4	42
34	Biosynthesis and Biotechnology of High-Value p-Menthane Monoterpenes, Including Menthol, Carvone, and Limonene. <i>Advances in Biochemical Engineering/Biotechnology</i> , 2015, 148, 319-353.	0.6	41
35	Metabolite profiling of Calvin cycle intermediates by HPLC-MS using mixed-mode stationary phases. <i>Plant Journal</i> , 2008, 55, 1047-1060.	2.8	38
36	Morphology of glandular trichomes of Japanese catnip ( <i>Schizonepeta tenuifolia</i> Briquet) and developmental dynamics of their secretory activity. <i>Phytochemistry</i> , 2018, 150, 23-30.	1.4	35

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37	Multiple Levels of Regulation Determine Monoterpenoid Essential Oil Compositional Variation in the Mint Family. <i>Molecular Plant</i> , 2015, 8, 188-191.	3.9	32
38	NMR spectroscopic search module for Spektraris, an online resource for plant natural product identification – Taxane diterpenoids from <i>Taxus</i> —media cell suspension cultures as a case study. <i>Phytochemistry</i> , 2015, 113, 87-95.	1.4	32
39	Single-cell genomics. <i>Current Opinion in Plant Biology</i> , 2005, 8, 236-241.	3.5	30
40	Counting the cost of a cold-blooded life: Metabolomics of cold acclimation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2004, 101, 14996-14997.	3.3	29
41	Comprehensive Assessment of Transcriptional Regulation Facilitates Metabolic Engineering of Isoprenoid Accumulation in <i>Arabidopsis</i> . <i>Plant Physiology</i> , 2015, 169, pp.00573.2015.	2.3	29
42	Biosynthesis of Diterpenoids in <i>Tripterygium</i> Adventitious Root Cultures. <i>Plant Physiology</i> , 2017, 175, 92-103.	2.3	27
43	Bioenergetics of Monoterpenoid Essential Oil Biosynthesis in Nonphotosynthetic Glandular Trichomes. <i>Plant Physiology</i> , 2017, 175, 681-695.	2.3	23
44	Assessment of flux through oleoresin biosynthesis in epithelial cells of loblolly pine resin ducts. <i>Journal of Experimental Botany</i> , 2019, 70, 217-230.	2.4	22
45	Crop Wild Relatives as Germplasm Resource for Cultivar Improvement in Mint ( <i>Mentha</i> L.). <i>Frontiers in Plant Science</i> , 2020, 11, 1217.	1.7	22
46	Genome-Wide Analysis of Terpene Synthase Gene Family in <i>Mentha longifolia</i> and Catalytic Activity Analysis of a Single Terpene Synthase. <i>Genes</i> , 2021, 12, 518.	1.0	22
47	Taxanes and taxoids of the genus <i>Taxus</i> – A comprehensive inventory of chemical diversity. <i>Phytochemistry</i> , 2021, 190, 112829.	1.4	22
48	Enzymology of monoterpene functionalization in glandular trichomes. <i>Journal of Experimental Botany</i> , 2019, 70, 1095-1108.	2.4	21
49	Assessing Flux Distribution Associated with Metabolic Specialization of Glandular Trichomes. <i>Trends in Plant Science</i> , 2018, 23, 638-647.	4.3	20
50	Flavonoid deficiency disrupts redox homeostasis and terpenoid biosynthesis in glandular trichomes of tomato. <i>Plant Physiology</i> , 2022, 188, 1450-1468.	2.3	20
51	Misexpression of the Niemann-Pick disease type C1 (NPC1)-like protein in <i>Arabidopsis</i> causes sphingolipid accumulation and reproductive defects. <i>Planta</i> , 2015, 242, 921-933.	1.6	19
52	Metabolic shifts associated with drought-induced senescence in <i>Brachypodium</i> . <i>Plant Science</i> , 2019, 289, 110278.	1.7	18
53	Assessing Chemical Diversity in <i>Psilotum nudum</i> (L.) Beauv., a Pantropical Whisk Fern That Has Lost Many of Its Fern-Like Characters. <i>Frontiers in Plant Science</i> , 2019, 10, 868.	1.7	14
54	Genetic diversity survey of <i>Mentha aquatica</i> L. and <i>Mentha suaveolens</i> Ehrh., mint crop ancestors. <i>Genetic Resources and Crop Evolution</i> , 2019, 66, 825-845.	0.8	14

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55	Biochemical characterization of acyl activating enzymes for side chain moieties of Taxol and its analogs. <i>Journal of Biological Chemistry</i> , 2020, 295, 4963-4973.	1.6	13
56	Integrative analysis of metabolic networks: from peaks to flux models?. <i>Current Opinion in Plant Biology</i> , 2006, 9, 220-226.	3.5	12
57	Experimental sink removal induces stress responses, including shifts in amino acid and phenylpropanoid metabolism, in soybean leaves. <i>Planta</i> , 2012, 235, 939-954.	1.6	12
58	Validation of a microscale extraction and high-throughput UHPLC-QTOF-MS analysis method for huperzine A in <i>Huperzia</i> . <i>Biomedical Chromatography</i> , 2012, 26, 1191-1195.	0.8	11
59	Determinants of Enantiospecificity in Limonene Synthases. <i>Biochemistry</i> , 2020, 59, 1661-1664.	1.2	11
60	<i>bHLH093/NFL</i> and <i>bHLH061</i> are required for apical meristem function in <i>Arabidopsis thaliana</i> . <i>Plant Signaling and Behavior</i> , 2018, 13, e1486146.	1.2	10
61	Ultrastructure of Grapefruit Secretory Cavities and Immunocytochemical Localization of (+)-Limonene Synthase. <i>International Journal of Plant Sciences</i> , 2015, 176, 643-661.	0.6	9
62	Rapid purification of gram quantities of $\beta$ -sitosterol from a commercial phytosterol mixture. <i>BMC Research Notes</i> , 2014, 7, 182.	0.6	8
63	Generation and Functional Evaluation of Designer Monoterpene Synthases. <i>Methods in Enzymology</i> , 2016, 576, 147-165.	0.4	8
64	Altering potato isoprenoid metabolism increases biomass and induces early flowering. <i>Journal of Experimental Botany</i> , 2020, 71, 4109-4124.	2.4	8
65	Comprehensive inventory of cannabinoids in <i>Cannabis sativa</i> L.: Can we connect genotype and chemotype?. <i>Phytochemistry Reviews</i> , 2022, 21, 1273-1313.	3.1	8
66	Chromosome-level genome assembly of <i>Mentha longifolia</i> L. reveals gene organization underlying disease resistance and essential oil traits. <i>G3: Genes, Genomes, Genetics</i> , 2022, 12, .	0.8	8
67	Commercial-Scale Tissue Culture for the Production of Plant Natural Products: Successes, Failures and Outlook. , 2018, , 189-218.		6
68	Determinants of Selectivity for the Formation of Monocyclic and Bicyclic Products in Monoterpene Synthases. <i>ACS Catalysis</i> , 2022, 12, 7453-7469.	5.5	6
69	Soybean vegetative lipoxygenases are not vacuolar storage proteins. <i>Functional Plant Biology</i> , 2011, 38, 778.	1.1	5
70	Kinetic Modeling of Plant Metabolism and Its Predictive Power: Peppermint Essential Oil Biosynthesis as an Example. <i>Methods in Molecular Biology</i> , 2014, 1083, 287-311.	0.4	5
71	Sample Preparation for Single Cell Transcriptomics: Essential Oil Glands in Citrus Fruit Peel as an Example. <i>Methods in Molecular Biology</i> , 2014, 1153, 203-212.	0.4	5
72	Functional Characterization and Structural Insights Into Stereoselectivity of Pulegone Reductase in Menthol Biosynthesis. <i>Frontiers in Plant Science</i> , 2021, 12, 780970.	1.7	5

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73	Cell Type-Specific Transcriptome Analysis of the Soybean Leaf Paraveinal Mesophyll Layer. <i>Plant Molecular Biology Reporter</i> , 2013, 31, 210-221.	1.0	4
74	Online resources for gene discovery and biochemical research with aromatic and medicinal plants. <i>Phytochemistry Reviews</i> , 2016, 15, 489-510.	3.1	4
75	National Academies report has broad support. <i>Nature Biotechnology</i> , 2017, 35, 304-306.	9.4	3
76	Differential Accumulation of Metabolites and Transcripts Related to Flavonoid, Styrylpyrone, and Galactolipid Biosynthesis in Equisetum Species and Tissue Types. <i>Metabolites</i> , 2022, 12, 403.	1.3	3
77	Selectivity of enzymes involved in the formation of opposite enantiomeric series of p-menthane monoterpenoids in peppermint and Japanese catnip. <i>Plant Science</i> , 2022, 314, 111119.	1.7	1
78	Chapter six Genomic survey of metabolic pathways in rice. <i>Recent Advances in Phytochemistry</i> , 2004, 38, 111-137.	0.5	0
79	Comprehensive Post-Genomic Data Analysis Approaches Integrating Biochemical Pathway Maps. <i>ChemInform</i> , 2005, 36, no.	0.1	0
80	Multiple levels of regulation determine monoterpenoid essential oil compositional variation in the mint family. <i>Molecular Plant</i> , 2014, , .	3.9	0