## Sang Un Park

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Treasure from garden: Bioactive compounds of buckwheat. Food Chemistry, 2021, 335, 127653.	4.2	117

2 Anthocyanin Accumulation and Expression of Anthocyanin Biosynthetic Genes in Radish (<i>Raphanus) Tj ETQq0 0.0 rgBT /Oyerlock 10

3	Differential Expression of Flavonoid Biosynthesis Genes and Accumulation of Phenolic Compounds in Common Buckwheat ( <i>Fagopyrum esculentum</i> ). Journal of Agricultural and Food Chemistry, 2010, 58, 12176-12181.	2.4	98
4	Molecular cloning and characterization of phenylalanine ammonia-lyase, cinnamate 4-hydroxylase and genes involved in flavone biosynthesis in Scutellaria baicalensis. Bioresource Technology, 2010, 101, 9715-9722.	4.8	88
5	An up-to-date review of rutin and its biological and pharmacological activities. EXCLI Journal, 2015, 14, 59-63.	0.5	85
6	Effect of Different Agrobacterium rhizogenes Strains on Hairy Root Induction and Phenylpropanoid Biosynthesis in Tartary Buckwheat (Fagopyrum tataricum Gaertn). Frontiers in Microbiology, 2016, 7, 318.	1.5	83
7	Effects of White, Blue, and Red Light-Emitting Diodes on Carotenoid Biosynthetic Gene Expression Levels and Carotenoid Accumulation in Sprouts of Tartary Buckwheat (Fagopyrum tataricum Gaertn.). Journal of Agricultural and Food Chemistry, 2013, 61, 12356-12361.	2.4	79
8	Effects of Light-Emitting Diodes on Expression of Phenylpropanoid Biosynthetic Genes and Accumulation of Phenylpropanoids in <i>Fagopyrum tataricum</i> Sprouts. Journal of Agricultural and Food Chemistry, 2014, 62, 4839-4845.	2.4	79
9	Transcriptome analysis and metabolic profiling of green and red kale (Brassica oleracea var. acephala) seedlings. Food Chemistry, 2018, 241, 7-13.	4.2	75
10	Metabolomics Analysis and Biosynthesis of Rosmarinic Acid in Agastache rugosa Kuntze Treated with Methyl Jasmonate. PLoS ONE, 2013, 8, e64199.	1.1	73
11	Anthocyanin and Carotenoid Contents in Different Cultivars of Chrysanthemum (Dendranthema) Tj ETQq1 1 0.7	84314 rgB 1.7	BT /Qverlock
11	Anthocyanin and Carotenoid Contents in Different Cultivars of Chrysanthemum (Dendranthema) Tj ETQq1 1 0.7 Metabolic Profiling of Glucosinolates, Anthocyanins, Carotenoids, and Other Secondary Metabolites in Kohlrabi ( <i>Brassica oleracea</i> var. <i>gongylodes</i> ). Journal of Agricultural and Food Chemistry, 2012, 60, 8111-8116.	84314 rgE 1.7 2.4	70 T
11 12 13	Anthocyanin and Carotenoid Contents in Different Cultivars of Chrysanthemum (Dendranthema) Tj ETQq1 1 0.7 Metabolic Profiling of Glucosinolates, Anthocyanins, Carotenoids, and Other Secondary Metabolites in Kohlrabi ( <i>Brassica oleracea</i> var. <i>gongylodes</i> ). Journal of Agricultural and Food Chemistry, 2012, 60, 8111-8116. MYB Transcription Factors Regulate Glucosinolate Biosynthesis in Different Organs of Chinese Cabbage (Brassica rapa ssp. pekinensis). Molecules, 2013, 18, 8682-8695.	84314 rgE 2.4 1.7	8T <u>/Q</u> verlock 70 68
11 12 13 14	Anthocyanin and Carotenoid Contents in Different Cultivars of Chrysanthemum (Dendranthema) Tj ETQq1 1 0.7         Metabolic Profiling of Glucosinolates, Anthocyanins, Carotenoids, and Other Secondary Metabolites         in Kohlrabi ( <i>Brassica oleracea</i> var. <i>gongylodes</i> ). Journal of Agricultural and Food         Chemistry, 2012, 60, 8111-8116.         MYB Transcription Factors Regulate Glucosinolate Biosynthesis in Different Organs of Chinese         Cabbage (Brassica rapa ssp. pekinensis). Molecules, 2013, 18, 8682-8695.         Overexpression of a tartary buckwheat R2R3-MYB transcription factor gene, FtMYB9, enhances         tolerance to drought and salt stresses in transgenic Arabidopsis. Journal of Plant Physiology, 2017, 214, 81-90.	84314 rgB 2.4 1.7 1.6	8T <u>/Q</u> verlock 70 68 68
11 12 13 14 15	Anthocyanin and Carotenoid Contents in Different Cultivars of Chrysanthemum (Dendranthema) Tj ETQq1 1 0.7         Metabolic Profiling of Glucosinolates, Anthocyanins, Carotenoids, and Other Secondary Metabolites in Kohlrabi ( <i>Brassica oleracea</i> var. <i>gongylodes</i> ). Journal of Agricultural and Food Chemistry, 2012, 60, 8111-8116.         MYB Transcription Factors Regulate Glucosinolate Biosynthesis in Different Organs of Chinese Cabbage (Brassica rapa ssp. pekinensis). Molecules, 2013, 18, 8682-8695.         Overexpression of a tartary buckwheat R2R3-MYB transcription factor gene, FtMYB9 , enhances tolerance to drought and salt stresses in transgenic Arabidopsis. Journal of Plant Physiology, 2017, 214, 81-90.         Variation of glucosinolates in 62 varieties of Chinese cabbage (Brassica rapa L. ssp. pekinensis) and their antioxidant activity. LWT - Food Science and Technology, 2014, 58, 93-101.	84314 rgE 2.4 1.7 1.6 2.5	8T /Qverlock 70 68 68 66
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<ol> <li>11</li> <li>12</li> <li>13</li> <li>14</li> <li>15</li> <li>16</li> <li>17</li> </ol>	Anthocyanin and Carotenoid Contents in Different Cultivars of Chrysanthemum (Dendranthema) Tj ETQq1 1 0.7         Metabolic Profiling of Glucosinolates, Anthocyanins, Carotenoids, and Other Secondary Metabolites in Kohlrabi ( <i>Brassica oleracea</i> var. <i>gongylodes</i> ). Journal of Agricultural and Food Chemistry, 2012, 60, 8111-8116.         MYB Transcription Factors Regulate Glucosinolate Biosynthesis in Different Organs of Chinese Cabbage (Brassica rapa ssp. pekinensis). Molecules, 2013, 18, 8682-8695.         Overexpression of a tartary buckwheat R2R3-MYB transcription factor gene, FtMYB9, enhances tolerance to drought and salt stresses in transgenic Arabidopsis. Journal of Plant Physiology, 2017, 214, 81-90.         Variation of glucosinolates in 62 varieties of Chinese cabbage (Brassica rapa L. ssp. pekinensis) and their antioxidant activity. LWT - Food Science and Technology, 2014, 58, 93-101.         Enhanced Triterpene Accumulation in <i>Panax ginseng</i> Hairy Roots Overexpressing Mevalonate-5-pyrophosphate Decarboxylase and Farnesyl Pyrophosphate Synthase. ACS Synthetic Biology, 2014, 3, 773-779.         Accumulation of Tilianin and Rosmarinic Acid and Expression of Phenylpropanoid Biosynthetic Genes in <i>Agastache rugosa</i>	84314 rgE 2.4 1.7 1.6 2.5 1.9 2.4	8T <u>/9</u> verlock 70 68 68 66 66 65

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19	Metabolic Profiling and Antioxidant Assay of Metabolites from Three Radish Cultivars (Raphanus) Tj ETQq1 1 0.784	1314 rgBT 1.7	/Qverlock
20	Carotenoid content and expression of phytoene synthase and phytoene desaturase genes in bitter melon (Momordica charantia). Food Chemistry, 2011, 126, 1686-1692.	4.2	56
21	Characterization of two tartary buckwheat <scp>R2R3â€MYB</scp> transcription factors and their regulation of proanthocyanidin biosynthesis. Physiologia Plantarum, 2014, 152, 431-440.	2.6	56
22	Metabolite Profiling Approach Reveals the Interface of Primary and Secondary Metabolism in Colored Cauliflowers (Brassica oleracea L. ssp. <i>botrytis</i> ). Journal of Agricultural and Food Chemistry, 2013, 61, 6999-7007.	2.4	55
23	Differential stress-response expression of two flavonol synthase genes and accumulation of flavonols in tartary buckwheat. Journal of Plant Physiology, 2013, 170, 1630-1636.	1.6	54
24	Differential Expression of Anthocyanin Biosynthetic Genes and Anthocyanin Accumulation in Tartary Buckwheat Cultivars â€~Hokkai T8' and â€~Hokkai T10'. Journal of Agricultural and Food Chemistry, 2011, 2356-2361.	5294	52
25	Accumulation of Phenylpropanoids and Correlated Gene Expression during the Development of Tartary Buckwheat Sprouts. Journal of Agricultural and Food Chemistry, 2012, 60, 5629-5635.	2.4	52
26	Enhanced Accumulation of Phytosterol and Triterpene in Hairy Root Cultures of Platycodon grandiflorum by Overexpression of Panax ginseng 3-Hydroxy-3-methylglutaryl-coenzyme A Reductase. Journal of Agricultural and Food Chemistry, 2013, 61, 1928-1934.	2.4	52
27	Growth and Rutin Production in Hairy Root Cultures of Buckwheat (Fagopyrum esculentumM.). Preparative Biochemistry and Biotechnology, 2007, 37, 239-246.	1.0	51
28	Enhancement of flavone levels through overexpression of chalcone isomerase in hairy root cultures of Scutellaria baicalensis. Functional and Integrative Genomics, 2011, 11, 491-496.	1.4	51
29	In Vitro Antioxidant and Antimicrobial Properties of Flower, Leaf, and Stem Extracts of Korean Mint. Antioxidants, 2019, 8, 75.	2.2	51
30	Rosmarinic acid production in hairy root cultures of Agastache rugosa Kuntze. World Journal of Microbiology and Biotechnology, 2008, 24, 969-972.	1.7	50
31	Accumulation of Anthocyanin and Associated Gene Expression in Radish Sprouts Exposed to Light and Methyl Jasmonate. Journal of Agricultural and Food Chemistry, 2013, 61, 4127-4132.	2.4	50
32	Herbicidal activity of formulated sorgoleone, a natural product of sorghum root exudate. Pest Management Science, 2014, 70, 252-257.	1.7	49
33	Yeast Extract and Silver Nitrate Induce the Expression of Phenylpropanoid Biosynthetic Genes and Induce the Accumulation of Rosmarinic Acid in Agastache rugosa Cell Culture. Molecules, 2016, 21, 426.	1.7	49
34	De novo transcriptome analysis and glucosinolate profiling in watercress (Nasturtium officinale R.) Tj ETQq0 0 0 rg	,BT /Overlc	›දፄ 10 Tf 50
35	Effects of Light-Emitting Diodes on the Accumulation of Glucosinolates and Phenolic Compounds in Sprouting Canola (Brassica napus L.). Foods, 2019, 8, 76.	1.9	49
36	Production of phenolic compounds in hairy root culture of tartary buckwheat (Fagopyrum tataricum) Tj ETQq0 0 C	rgBT /Ov€	erlock 10 Tf

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#	Article	IF	CITATIONS
37	Ginsenoside content of berries and roots of three typical Korean ginseng (Panax ginseng) cultivars. Natural Product Communications, 2009, 4, 903-6.	0.2	47
38	Effects of cold stress on transcripts and metabolites in tartary buckwheat (Fagopyrum tataricum). Environmental and Experimental Botany, 2018, 155, 488-496.	2.0	43
39	Identification of phenylpropanoid biosynthetic genes and phenylpropanoid accumulation by transcriptome analysis of Lycium chinense. BMC Genomics, 2013, 14, 802.	1.2	42
40	Accumulation of Carotenoids and Metabolic Profiling in Different Cultivars of Tagetes Flowers. Molecules, 2017, 22, 313.	1.7	42
41	Transcriptome Analysis and Metabolic Profiling of Lycoris Radiata. Biology, 2019, 8, 63.	1.3	42
42	Accumulation of Phenylpropanoids by White, Blue, and Red Light Irradiation and Their Organ-Specific Distribution in Chinese Cabbage ( <i>Brassica rapa</i> ssp. <i>pekinensis</i> ). Journal of Agricultural and Food Chemistry, 2015, 63, 6772-6778.	2.4	41
43	Metabolic Profiling in Chinese Cabbage ( <i>Brassica rapa</i> L. subsp. <i>pekinensis</i> ) Cultivars Reveals that Glucosinolate Content Is Correlated with Carotenoid Content. Journal of Agricultural and Food Chemistry, 2016, 64, 4426-4434.	2.4	41
44	Influence of Indole-3-Acetic Acid and Gibberellic Acid on Phenylpropanoid Accumulation in Common Buckwheat (Fagopyrum esculentum Moench) Sprouts. Molecules, 2017, 22, 374.	1.7	40
45	Enhancement of Anti-Inflammatory Activity of Aloe vera Adventitious Root Extracts through the Alteration of Primary and Secondary Metabolites via Salicylic Acid Elicitation. PLoS ONE, 2013, 8, e82479.	1.1	40
46	An update on biosynthesis and regulation of carotenoids in plants. South African Journal of Botany, 2021, 140, 290-302.	1.2	39
47	An efficient protocol for genetic transformation of watercress (Nasturtium officinale) using Agrobacterium rhizogenes. Molecular Biology Reports, 2011, 38, 4947-4953.	1.0	38
48	Metabolomic Analysis and Phenylpropanoid Biosynthesis in Hairy Root Culture of Tartary Buckwheat Cultivars. PLoS ONE, 2013, 8, e65349.	1.1	38
49	Accumulation of Rutin and Betulinic Acid and Expression of Phenylpropanoid and Triterpenoid Biosynthetic Genes in Mulberry ( <i>Morus alba</i> L.). Journal of Agricultural and Food Chemistry, 2015, 63, 8622-8630.	2.4	38
50	Physiological Roles of Rutin in the Buckwheat Plant. Japan Agricultural Research Quarterly, 2015, 49, 37-43.	0.1	37
51	Identification, isolation and expression analysis of eight stress-related R2R3-MYB genes in tartary buckwheat (Fagopyrum tataricum). Plant Cell Reports, 2016, 35, 1385-1396.	2.8	37
52	Influence of Chitosan, Salicylic Acid and Jasmonic Acid on Phenylpropanoid Accumulation in Germinated Buckwheat (Fagopyrum esculentum Moench). Foods, 2019, 8, 153.	1.9	37
53	Triterpene and Flavonoid Biosynthesis and Metabolic Profiling of Hairy Roots, Adventitious Roots, and Seedling Roots of <i>Astragalus membranaceus</i> . Journal of Agricultural and Food Chemistry, 2015, 63, 8862-8869.	2.4	36
54	Enhancement of rutin in Fagopyrum esculentum hairy root cultures by the Arabidopsis transcription factor AtMYB12. Biotechnology Letters, 2012, 34, 577-583.	1.1	35

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55	Metabolic Profiling of Nine Mentha Species and Prediction of Their Antioxidant Properties Using Chemometrics. Molecules, 2019, 24, 258.	1.7	35
56	Enhancement of the flavone contents of Scutellaria baicalensis hairy roots via metabolic engineering using maize Lc and Arabidopsis PAP1 transcription factors. Metabolic Engineering, 2021, 64, 64-73.	3.6	35
57	LED Lights Enhance Metabolites and Antioxidants in Chinese Cabbage and Kale. Brazilian Archives of Biology and Technology, 2016, 59, .	0.5	34
58	Phenolic compound profiles and their seasonal variations in new red-phenotype head-forming Chinese cabbages. LWT - Food Science and Technology, 2018, 90, 433-439.	2.5	34
59	Comparative Analysis of Flavonoids and Polar Metabolite Profiling of Tanno-Original and Tanno-High Rutin Buckwheat. Journal of Agricultural and Food Chemistry, 2014, 62, 2701-2708.	2.4	32
60	Molecular Cloning and Characterization of Phenylalanine Ammonia-lyase and Cinnamate 4-Hydroxylase in the Phenylpropanoid Biosynthesis Pathway in Garlic (Allium sativum). Journal of Agricultural and Food Chemistry, 2010, 58, 10911-10917.	2.4	31
61	Differentiated cuticular wax content and expression patterns of cuticular wax biosynthetic genes in bloomed and bloomless broccoli (Brassica oleracea var. italica). Process Biochemistry, 2015, 50, 456-462.	1.8	31
62	Metabolic profiling of pale green and purple kohlrabi (Brassica oleracea var. gongylodes). Applied Biological Chemistry, 2017, 60, 249-257.	0.7	31
63	Analysis of Metabolites in White Flowers of Magnolia Denudata Desr. and Violet Flowers of Magnolia Liliiflora Desr Molecules, 2018, 23, 1558.	1.7	31
64	Enhancing Sorgoleone Levels in Grain Sorghum Root Exudates. Journal of Chemical Ecology, 2010, 36, 914-922.	0.9	30
65	Comparative Metabolic Profiling of Green and Purple Pakchoi (Brassica Rapa Subsp. Chinensis). Molecules, 2018, 23, 1613.	1.7	30
66	Effects of LED lights on Expression of Genes Involved in Phenylpropanoid Biosynthesis and Accumulation of Phenylpropanoids in Wheat Sprout. Agronomy, 2019, 9, 307.	1.3	30
67	Metabolite Profiling and Comparative Analysis of Secondary Metabolites in Chinese Cabbage, Radish, and Hybrid <i>xBrassicoraphanus</i> . Journal of Agricultural and Food Chemistry, 2020, 68, 13711-13719.	2.4	30
68	Molecular characterisation and the light–dark regulation of carotenoid biosynthesis in sprouts of tartary buckwheat (Fagopyrum tataricum Gaertn.). Food Chemistry, 2013, 141, 3803-3812.	4.2	29
69	Influence of Light on the Free Amino Acid Content and γ-Aminobutyric Acid Synthesis in Brassica juncea Seedlings. Journal of Agricultural and Food Chemistry, 2013, 61, 8624-8631.	2.4	29
70	Metabolic Differentiation of Diamondback Moth (Plutella xylostella (L.)) Resistance in Cabbage (Brassica oleracea L. ssp. <i>capitata</i> ). Journal of Agricultural and Food Chemistry, 2013, 61, 11222-11230.	2.4	28
71	Identification and quantification of carotenoids in paprika fruits and cabbage, kale, and lettuce leaves. Journal of the Korean Society for Applied Biological Chemistry, 2014, 57, 355-358.	0.9	28
72	Far infrared irradiation alters total polyphenol, total flavonoid, antioxidant property and quercetin production in tartary buckwheat sprout powder. Journal of Cereal Science, 2014, 59, 167-172.	1.8	28

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73	Medically important carotenoids from Momordica charantia and their gene expressions in different organs. Saudi Journal of Biological Sciences, 2017, 24, 1913-1919.	1.8	28
74	Molecular characterization of anthocyanin and betulinic acid biosynthesis in red and white mulberry fruits using high-throughput sequencing. Food Chemistry, 2019, 279, 364-372.	4.2	28
75	Effect of Salinity Stress on Phenylpropanoid Genes Expression and Related Gene Expression in Wheat Sprout. Agronomy, 2020, 10, 390.	1.3	28
76	Comparative Analysis of Secondary Metabolites and Metabolic Profiling between Diploid and Tetraploid <i>Morus alba</i> L Journal of Agricultural and Food Chemistry, 2021, 69, 1300-1307.	2.4	28
77	Metabolomic Analysis and Differential Expression of Anthocyanin Biosynthetic Genes in White- and Red-Flowered Buckwheat Cultivars (Fagopyrum esculentum). Journal of Agricultural and Food Chemistry, 2013, 61, 10525-10533.	2.4	27
78	Enhancement of Chlorogenic Acid Production in Hairy Roots of Platycodon grandiflorum by Over-Expression of An Arabidopsis thaliana Transcription Factor AtPAP1. International Journal of Molecular Sciences, 2014, 15, 14743-14752.	1.8	27
79	Comparative analysis of flavonoids and polar metabolites from hairy roots of Scutellaria baicalensis and Scutellaria lateriflora. World Journal of Microbiology and Biotechnology, 2014, 30, 887-892.	1.7	27
80	Metabolic profiling and antioxidant activity during flower development in Agastache rugosa. Physiology and Molecular Biology of Plants, 2021, 27, 445-455.	1.4	27
81	Influence of light-emitting diodes on phenylpropanoid biosynthetic gene expression and phenylpropanoid accumulation in Agastache rugosa. Applied Biological Chemistry, 2020, 63, .	0.7	27
82	Age-dependent Distribution of Fungal Endophytes in Panax ginseng Roots Cultivated in Korea. Journal of Ginseng Research, 2012, 36, 327-333.	3.0	27
83	Comparative Phytochemical Analyses and Metabolic Profiling of Different Phenotypes of Chinese Cabbage (Brassica Rapa ssp. Pekinensis). Foods, 2019, 8, 587.	1.9	26
84	Cloning and Characterization of Phenylalanine Ammonia-Lyase and Cinnamate 4-Hydroxylase and Pyranocoumarin Biosynthesis in <i>Angelica gigas</i> . Journal of Natural Products, 2010, 73, 1394-1397.	1.5	25
85	Overexpression of phenylalanine ammonia-lyase improves flavones production in transgenic hairy root cultures of Scutellaria baicalensis. Process Biochemistry, 2012, 47, 2575-2580.	1.8	25
86	Metabolomics for the Quality Assessment of <i>Lycium chinense</i> Fruits. Bioscience, Biotechnology and Biochemistry, 2012, 76, 2188-2194.	0.6	24
87	Regulation of the major vacuolar Ca2+ transporter genes, by intercellular Ca2+ concentration and abiotic stresses, in tip-burn resistant Brassica oleracea. Molecular Biology Reports, 2013, 40, 177-188.	1.0	24
88	Determination of lipophilic metabolites for species discrimination and quality assessment of nine leafy vegetables. Journal of the Korean Society for Applied Biological Chemistry, 2015, 58, 909-918.	0.9	24
89	Fungal Endophytes from Three Cultivars of Panax ginseng Meyer Cultivated in Korea. Journal of Ginseng Research, 2012, 36, 107-113.	3.0	23
90	Effects of Light-Emitting Diodes on the Accumulation of Phenolic Compounds and Glucosinolates in Brassica juncea Sprouts. Horticulturae, 2020, 6, 77.	1.2	23

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91	Accumulation of Anthocyanins through Overexpression of AtPAP1 in Solanum nigrum Lin. (Black) Tj ETQq1 1 0.78	4314 rgB1 1.8	7 / Gverlock
92	Biological activity of various radish species. Oriental Pharmacy and Experimental Medicine, 2015, 15, 105-111.	1.2	22
93	Molecular cloning and characterization of rosmarinic acid biosynthetic genes and rosmarinic acid accumulation in Ocimum basilicum L. Saudi Journal of Biological Sciences, 2019, 26, 469-472.	1.8	22
94	Metabolic Profiling-Based Evaluation of the Fermentative Behavior of Aspergillus oryzae and Bacillus subtilis for Soybean Residues Treated at Different Temperatures. Foods, 2020, 9, 117.	1.9	22
95	Recent studies on rosmarinic acid and its biological and pharmacological activities. EXCLI Journal, 2014, 13, 1192-5.	0.5	22
96	Effects of Jasmonates on Sorgoleone Accumulation and Expression of Genes for Sorgoleone Biosynthesis in Sorghum Roots. Journal of Chemical Ecology, 2013, 39, 712-722.	0.9	21
97	Red Chinese Cabbage Transcriptome Analysis Reveals Structural Genes and Multiple Transcription Factors Regulating Reddish Purple Color. International Journal of Molecular Sciences, 2020, 21, 2901.	1.8	21
98	Yeast extract improved biosynthesis of astragalosides in hairy root cultures of <i>Astragalus membranaceus</i> . Preparative Biochemistry and Biotechnology, 2021, 51, 467-474.	1.0	21
99	Metabolomic analysis reveals the interaction of primary and secondary metabolism in white, pale green, and green pak choi (Brassica rapa subsp. chinensis). Applied Biological Chemistry, 2021, 64, .	0.7	21
100	Profiles of Secondary Metabolites (Phenolic Acids, Carotenoids, Anthocyanins, and Galantamine) and Primary Metabolites (Carbohydrates, Amino Acids, and Organic Acids) during Flower Development in Lycoris radiata. Biomolecules, 2021, 11, 248.	1.8	21
101	Transcriptional Profiling and Molecular Characterization of Astragalosides, Calycosin, and Calycosin-7- <i>O</i> -î <sup>2</sup> - <scp>d</scp> -glucoside Biosynthesis in the Hairy Roots of <i>Astragalus membranaceus</i> in Response to Methyl Jasmonate. Journal of Agricultural and Food Chemistry, 2015, 63 6231-6240	2.4	20
102	A Comparative Study of Phenolic Antioxidant Activity and Flavonoid Biosynthesisâ€Related Gene Expression Between Summer and Winter Strawberry Cultivars. Journal of Food Science, 2017, 82, 341-349.	1.5	20
103	Metabolomic Profiling of the White, Violet, and Red Flowers of Rhododendron schlippenbachii Maxim Molecules, 2018, 23, 827.	1.7	20
104	Accumulation of Flavonoids and Related Gene Expressions in Different Organs of Astragalus membranaceus Bge. Applied Biochemistry and Biotechnology, 2014, 173, 2076-2085.	1.4	19
105	Molecular Cloning and Characterization of Genes Involved in Rosmarinic Acid Biosynthesis from <i>Prunella vulgaris</i> . Biological and Pharmaceutical Bulletin, 2014, 37, 1221-1227.	0.6	19
106	Fagopyrum tataricum FtWD40 Functions as a Positive Regulator of Anthocyanin Biosynthesis in Transgenic Tobacco. Journal of Plant Growth Regulation, 2017, 36, 755-765.	2.8	19
107	Molecular characterization of flavonoid biosynthetic genes and accumulation of baicalin, baicalein, and wogonin in plant and hairy root of Scutellaria lateriflora. Saudi Journal of Biological Sciences, 2018, 25, 1639-1647.	1.8	19
108	Carotenoid Accumulation and Characterization of cDNAs Encoding Phytoene Synthase and Phytoene Desaturase in Garlic ( <i>Allium sativum</i> ). Journal of Agricultural and Food Chemistry, 2011, 59, 5412-5417.	2.4	18

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109	Overexpression of Cinnamate 4-Hydroxylase Gene Enhances Biosynthesis of Decursinol Angelate in Angelica gigas Hairy Roots. Molecular Biotechnology, 2012, 50, 114-120.	1.3	18
110	Characterization of Genes for a Putative Hydroxycinnamoyl-coenzyme A Quinate Transferase and <i>p</i> -Coumarate 3′-Hydroxylase and Chlorogenic Acid Accumulation in Tartary Buckwheat. Journal of Agricultural and Food Chemistry, 2013, 61, 4120-4126.	2.4	18
111	Accumulation of Phenylpropanoids and Correlated Gene Expression in Hairy Roots of Tartary Buckwheat under Light and Dark Conditions. Applied Biochemistry and Biotechnology, 2014, 174, 2537-2547.	1.4	18
112	Ginseng: a miracle sources of herbal and pharmacological uses. Oriental Pharmacy and Experimental Medicine, 2016, 16, 243-250.	1.2	18
113	Accumulation of Charantin and Expression of Triterpenoid Biosynthesis Genes in Bitter Melon ( <i>Momordica charantia</i> ). Journal of Agricultural and Food Chemistry, 2017, 65, 7240-7249.	2.4	18
114	Transcriptome Analysis and Metabolic Profiling of Green and Red Mizuna (Brassica rapa L. var.) Tj ETQq0 0 0 rg	BT /Qyerloc	k 10 Tf 50 54
115	Production of rosmarinic acid and correlated gene expression in hairy root cultures of green and purple basil ( <i>Ocimum basilicum</i> L.). Preparative Biochemistry and Biotechnology, 2021, 51, 35-43.	1.0	18
116	Metabolite Profiling Reveals Distinct Modulation of Complex Metabolic Networks in Non-Pigmented, Black, and Red Rice (Oryza sativa L.) Cultivars. Metabolites, 2021, 11, 367.	1.3	18
117	Molecular Characterization of Carotenoid Cleavage Dioxygenases and the Effect of Gibberellin, Abscisic Acid, and Sodium Chloride on the Expression of Genes Involved in the Carotenoid Biosynthetic Pathway and Carotenoid Accumulation in the Callus of Scutellaria baicalensis Georgi. Journal of Agricultural and Food Chemistry, 2013, 61, 5565-5572.	2.4	17
118	Accumulation of Kaempferitrin and Expression of Phenyl-Propanoid Biosynthetic Genes in Kenaf (Hibiscus cannabinus). Molecules, 2014, 19, 16987-16997.	1.7	17
119	Accumulation of Astragalosides and Related Gene Expression in Different Organs of Astragalus Membranaceus Bge. var Mongholicus (Bge.). Molecules, 2014, 19, 10922-10935.	1.7	17
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