Alisdair R Fernie

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

 596
 42,481
 109
 186

 papers
 citations
 h-index
 g-index

 704
 53,091
 8.7
 7.82

704 ext. papers

53,091 ext. citations

avg, IF

7.82 L-index

#	Paper	IF	Citations
596	The reliance of phytohormone biosynthesis on primary metabolite precursors <i>Journal of Plant Physiology</i> , 2022 , 268, 153589	3.6	1
595	The AtMYB60 transcription factor regulates stomatal opening by modulating oxylipin synthesis in guard cells <i>Scientific Reports</i> , 2022 , 12, 533	4.9	3
594	Unravelling the molecular networks that regulate kiwifruit flavor. <i>New Phytologist</i> , 2022 , 233, 8-10	9.8	
593	Environmentally-driven metabolite and lipid variations correspond to altered bioactivities of black wolfberry fruit. <i>Food Chemistry</i> , 2022 , 372, 131342	8.5	3
592	A Comparative Study of the Antihypertensive and Cardioprotective Potentials of Hot and Cold Aqueous Extracts of L. in Relation to Their Metabolic Profiles <i>Frontiers in Pharmacology</i> , 2022 , 13, 8404	1 78	1
591	Potential Valorization of Edible Nuts By-Products: Exploring the Immune-Modulatory and Antioxidants Effects of Selected Nut Shells Extracts in Relation to Their Metabolic Profiles <i>Antioxidants</i> , 2022 , 11,	7.1	9
590	Towards the Development, Maintenance and Standardized Phenotypic Characterization of Single-Seed-Descent Genetic Resources for Chickpea <i>Current Protocols</i> , 2022 , 2, e371		О
589	Plant metabolic gene clusters in the multi-omics era <i>Trends in Plant Science</i> , 2022 ,	13.1	3
588	Convergent selection of a WD40 protein that enhances grain yield in maize and rice <i>Science</i> , 2022 , 375, eabg7985	33.3	4
587	Auxin boosts energy generation pathways to fuel pollen maturation in barley <i>Current Biology</i> , 2022 ,	6.3	1
586	Maize Field Study Reveals Covaried Microbiota and Metabolic Changes in Roots over Plant Growth <i>MBio</i> , 2022 , e0258421	7.8	O
585	A reactive oxygen species burst causes haploid induction in maize Molecular Plant, 2022,	14.4	1
584	Bringing more players into play: Leveraging stress in genome wide association studies <i>Journal of Plant Physiology</i> , 2022 , 271, 153657	3.6	1
583	Natural variance at the interface of plant primary and specialized metabolism <i>Current Opinion in Plant Biology</i> , 2022 , 67, 102201	9.9	3
582	Rising rates of starch degradation during daytime and trehalose 6-phosphate optimize carbon availability <i>Plant Physiology</i> , 2022 ,	6.6	1
581	Measurement of Flower Metabolite Concentrations Using Gas Chromatography-Mass Spectrometry and High-Performance Liquid Chromatography-Mass Spectrometry <i>Methods in Molecular Biology</i> , 2022 , 2484, 3-12	1.4	
580	Preparation and Curation of Omics Data for Genome-Wide Association Studies. <i>Methods in Molecular Biology</i> , 2022 , 127-150	1.4	

579	Metabolomics Approaches for Studying the Trichoderma-Plant Interactions. <i>Rhizosphere Biology</i> , 2022 , 135-154	0.8	
578	Stable and Temporary Enzyme Complexes and Metabolons Involved in Energy and Redox Metabolism. <i>Antioxidants and Redox Signaling</i> , 2021 , 35, 788-807	8.4	7
577	The interface of central metabolism with hormone signaling in plants. Current Biology, 2021, 31, R1535	5-R61\$48	3 4
576	Enhancing crop diversity for food security in the face of climate uncertainty. Plant Journal, 2021,	6.9	9
575	The Arabidopsis electron-transfer flavoprotein: ubiquinone oxidoreductase is required during normal seed development and germination. <i>Plant Journal</i> , 2021 ,	6.9	1
574	High-quality reference genome sequences of two coconut cultivars provide insights into evolution of monocot chromosomes and differentiation of fiber content and plant height. <i>Genome Biology</i> , 2021 , 22, 304	18.3	3
573	Different Metabolic Roles for Alternative Oxidase in Leaves of Palustrine and Terrestrial Species. <i>Frontiers in Plant Science</i> , 2021 , 12, 752795	6.2	
572	Genetic variation in YIGE1 contributes to ear length and grain yield in maize. New Phytologist, 2021,	9.8	2
571	Metabolism-mediated mechanisms underpin the differential stomatal speediness regulation among ferns and angiosperms. <i>Plant, Cell and Environment,</i> 2021 ,	8.4	1
570	Pathways to de novo domestication of crop wild relatives. <i>Plant Physiology</i> , 2021 ,	6.6	4
569	Genome-wide association studies of Arabidopsis dark-induced senescence reveals signatures of autophagy in metabolic reprogramming. <i>Autophagy</i> , 2021 , 1-2	10.2	1
568	Diversity: current and prospective secondary metabolites for nutrition and medicine <i>Current Opinion in Biotechnology</i> , 2021 , 74, 164-170	11.4	2
567	Metabolic shifts during fruit development in pungent and non-pungent peppers <i>Food Chemistry</i> , 2021 , 375, 131850	8.5	0
566	Genome-wide association of the metabolic shifts underpinning dark-induced senescence in Arabidopsis. <i>Plant Cell</i> , 2021 ,	11.6	6
565	Rice metabolic regulatory network spanning the entire life cycle. <i>Molecular Plant</i> , 2021 ,	14.4	5
564	Multi-omics approaches explain the growth-promoting effect of the apocarotenoid growth regulator zaxinone in rice. <i>Communications Biology</i> , 2021 , 4, 1222	6.7	3
563	The nutritional profile and human health benefit of pigmented rice and the impact of post-harvest processes and product development on the nutritional components: A review. <i>Critical Reviews in Food Science and Nutrition</i> , 2021 , 1-28	11.5	1
562	Pod indehiscence in common bean is associated with the fine regulation of PvMYB26. <i>Journal of Experimental Botany</i> , 2021 , 72, 1617-1633	7	3

561	The cytosolic invertase NI6 affects vegetative growth, flowering, fruit set, and yield in tomato. Journal of Experimental Botany, 2021 , 72, 2525-2543	7	2
560	Multi-omics approach reveals the contribution of KLU to leaf longevity and drought tolerance. <i>Plant Physiology</i> , 2021 , 185, 352-368	6.6	8
559	Kingdom-wide analysis of the evolution of the plant type III polyketide synthase superfamily. <i>Plant Physiology</i> , 2021 , 185, 857-875	6.6	5
558	Cross-Species Metabolic Profiling of Floral Specialized Metabolism Facilitates Understanding of Evolutional Aspects of Metabolism Among Brassicaceae Species. <i>Frontiers in Plant Science</i> , 2021 , 12, 64	0 ⁶ 2 1	1
557	Several geranylgeranyl diphosphate synthase isoforms supply metabolic substrates for carotenoid biosynthesis in tomato. <i>New Phytologist</i> , 2021 , 231, 255-272	9.8	14
556	The genetics underlying metabolic signatures in a brown rice diversity panel and their vital role in human nutrition. <i>Plant Journal</i> , 2021 , 106, 507-525	6.9	6
555	Meeting human dietary vitamin requirements in the staple rice via strategies of biofortification and post-harvest fortification. <i>Trends in Food Science and Technology</i> , 2021 , 109, 65-82	15.3	9
554	Downregulation of the E2 Subunit of 2-Oxoglutarate Dehydrogenase Modulates Plant Growth by Impacting Carbon-Nitrogen Metabolism in Arabidopsis thaliana. <i>Plant and Cell Physiology</i> , 2021 , 62, 798	s- 8 194	3
553	Using landrace transcription factor alleles to increase yield in modern rice under low input agriculture. <i>Journal of Plant Physiology</i> , 2021 , 258-259, 153362	3.6	1
552	Multiomics-based dissection of citrus flavonoid metabolism using a Citrus reticulata iPoncirus trifoliata population. <i>Horticulture Research</i> , 2021 , 8, 56	7.7	11
551	Multi-omics analysis of early leaf development in. <i>Patterns</i> , 2021 , 2, 100235	5.1	4
550	Using precision phenotyping to inform de novo domestication. <i>Plant Physiology</i> , 2021 , 186, 1397-1411	6.6	1
549	Mild reductions in guard cell sucrose synthase 2 expression leads to slower stomatal opening and decreased whole plant transpiration in Nicotiana tabacum L. <i>Environmental and Experimental Botany</i> , 2021 , 184, 104370	5.9	2
548	Diversity of Chemical Structures and Biosynthesis of Polyphenols in Nut-Bearing Species. <i>Frontiers in Plant Science</i> , 2021 , 12, 642581	6.2	3
547	Associating primary and specialized metabolism with salt induced osmotic stress tolerance in maize. <i>New Phytologist</i> , 2021 , 230, 2091-2093	9.8	1
546	Differential responses of three Urochloa species to low phosphorus availability. <i>Annals of Applied Biology</i> , 2021 , 179, 216-230	2.6	O
545	From flowers to seeds: how the metabolism of flowers frames plant reproduction. <i>Biochemist</i> , 2021 , 43, 14-18	0.5	
544	Phosphoglycerate dehydrogenase genes differentially affect Arabidopsis metabolism and development. <i>Plant Science</i> , 2021 , 306, 110863	5.3	1

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543	The phosphorylated pathway of serine biosynthesis links plant growth with nitrogen metabolism. <i>Plant Physiology</i> , 2021 , 186, 1487-1506	6.6	0
542	Towards the Development, Maintenance, and Standardized Phenotypic Characterization of Single-Seed-Descent Genetic Resources for Common Bean. <i>Current Protocols</i> , 2021 , 1, e133		5
541	Overexpression of thioredoxin m in chloroplasts alters carbon and nitrogen partitioning in tobacco. Journal of Experimental Botany, 2021 , 72, 4949-4964	7	0
540	Ultra-high-performance liquid chromatography high-resolution mass spectrometry variants for metabolomics research. <i>Nature Methods</i> , 2021 , 18, 733-746	21.6	28
539	CsbZIP1-CsMYB12 mediates the production of bitter-tasting flavonols in tea plants (Camellia sinensis) through a coordinated activator-repressor network. <i>Horticulture Research</i> , 2021 , 8, 110	7.7	16
538	Metabolite Profiling in with Moderately Impaired Photorespiration Reveals Novel Metabolic Links and Compensatory Mechanisms of Photorespiration. <i>Metabolites</i> , 2021 , 11,	5.6	2
537	Domestication of Crop Metabolomes: Desired and Unintended Consequences. <i>Trends in Plant Science</i> , 2021 , 26, 650-661	13.1	14
536	When a Crop Goes Back to the Wild: Feralization. <i>Trends in Plant Science</i> , 2021 , 26, 543-545	13.1	4
535	Tyr-Asp inhibition of glyceraldehyde 3-phosphate dehydrogenase affects plant redox metabolism. <i>EMBO Journal</i> , 2021 , 40, e106800	13	8
534	Multiomics analyses reveal the roles of the ASR1 transcription factor in tomato fruits. <i>Journal of Experimental Botany</i> , 2021 , 72, 6490-6509	7	2
533	Two mitochondrial phosphatases, PP2c63 and Sal2, are required for posttranslational regulation of the TCA cycle in Arabidopsis. <i>Molecular Plant</i> , 2021 , 14, 1104-1118	14.4	9
532	Mass spectrometry-based metabolomics: a guide for annotation, quantification and best reporting practices. <i>Nature Methods</i> , 2021 , 18, 747-756	21.6	83
531	Exploring the genic resources underlying metabolites through mGWAS and mQTL in wheat: From large-scale gene identification and pathway elucidation to crop improvement. <i>Plant Communications</i> , 2021 , 2, 100216	9	2
530	Plasticity of rosette size in response to nitrogen availability is controlled by an RCC1-family protein. <i>Plant, Cell and Environment</i> , 2021 , 44, 3398-3411	8.4	2
529	Crop breeding - From experience-based selection to precision design. <i>Journal of Plant Physiology</i> , 2021 , 256, 153313	3.6	4
528	Auto-deconvolution and molecular networking of gas chromatography-mass spectrometry data. <i>Nature Biotechnology</i> , 2021 , 39, 169-173	44.5	36
527	Metabolons, enzyme-enzyme assemblies that mediate substrate channeling, and their roles in plant metabolism. <i>Plant Communications</i> , 2021 , 2, 100081	9	29
526	SWATH-MS-Based Proteomics: Strategies and Applications in Plants. <i>Trends in Biotechnology</i> , 2021 , 39, 433-437	15.1	18

525	Long-distance stress and developmental signals associated with abscisic acid signaling in environmental responses. <i>Plant Journal</i> , 2021 , 105, 477-488	6.9	11
524	Validated MAGIC and GWAS population mapping reveals the link between vitamin E content and natural variation in chorismate metabolism in tomato. <i>Plant Journal</i> , 2021 , 105, 907-923	6.9	4
523	Integrating multi-omics data for crop improvement. Journal of Plant Physiology, 2021, 257, 153352	3.6	24
522	A phased genome based on single sperm sequencing reveals crossover pattern and complex relatedness in tea plants. <i>Plant Journal</i> , 2021 , 105, 197-208	6.9	2
521	Decreased Levels of Thioredoxin Influences Stomatal Development and Aperture but Not Photosynthesis under Non-Stress and Saline Conditions. <i>International Journal of Molecular Sciences</i> , 2021 , 22,	6.3	3
520	Global mapping of protein-metabolite interactions in Saccharomyces cerevisiae reveals that Ser-Leu dipeptide regulates phosphoglycerate kinase activity. <i>Communications Biology</i> , 2021 , 4, 181	6.7	12
519	Evolutionary gain of oligosaccharide hydrolysis and sugar transport enhanced carbohydrate partitioning in sweet watermelon fruits. <i>Plant Cell</i> , 2021 , 33, 1554-1573	11.6	16
518	The NAC transcription factor FaRIF controls fruit ripening in strawberry. <i>Plant Cell</i> , 2021 , 33, 1574-1593	11.6	14
517	Thioredoxin-mediated regulation of (photo)respiration and central metabolism. <i>Journal of Experimental Botany</i> , 2021 , 72, 5987-6002	7	3
516	Phytochromes control metabolic flux, and their action at the seedling stage determines adult plant biomass. <i>Journal of Experimental Botany</i> , 2021 , 72, 3263-3278	7	1
515	The utility of metabolomics as a tool to inform maize biology. <i>Plant Communications</i> , 2021 , 2, 100187	9	3
514	Genome-wide association studies: assessing trait characteristics in model and crop plants. <i>Cellular and Molecular Life Sciences</i> , 2021 , 78, 5743-5754	10.3	10
513	Towards Development, Maintenance, and Standardized Phenotypic Characterization of Single-Seed-Descent Genetic Resources for Lupins. <i>Current Protocols</i> , 2021 , 1, e191		4
512	From Affinity to Proximity Techniques to Investigate Protein Complexes in Plants. <i>International Journal of Molecular Sciences</i> , 2021 , 22,	6.3	2
511	OsGF14b modulates defense signaling pathways in rice panicle blast response. <i>Crop Journal</i> , 2021 , 9, 725-738	4.6	5
510	The genomes of Taxus species unveil novel candidates in the biosynthesis of taxoids. <i>Molecular Plant</i> , 2021 , 14, 1773-1775	14.4	
509	Sulfur deficiency-induced genes affect seed protein accumulation and composition under sulfate deprivation. <i>Plant Physiology</i> , 2021 , 187, 2419-2434	6.6	5
508	Plant cell cultures as heterologous bio-factories for secondary metabolite production. <i>Plant Communications</i> , 2021 , 2, 100235	9	7

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507	Establishment of a GC-MS-based C-positional isotopomer approach suitable for investigating metabolic fluxes in plant primary metabolism. <i>Plant Journal</i> , 2021 , 108, 1213-1233	6.9	2
506	Genomic basis underlying the metabolome-mediated drought adaptation of maize. <i>Genome Biology</i> , 2021 , 22, 260	18.3	3
505	Plant metabolism paves the way for breeding crops with high nutritional value and stable yield. <i>Science China Life Sciences</i> , 2021 , 1	8.5	1
504	Will Stress Resilience Be Maintained in the Face of Climate Change?. <i>Metabolites</i> , 2021 , 11,	5.6	2
503	The INCREASE project: Intelligent Collections of food-legume genetic resources for European agrofood systems. <i>Plant Journal</i> , 2021 , 108, 646-660	6.9	5
502	The integration of MS-based metabolomics and multivariate data analysis allows for improved quality assessment of Zingiber officinale Roscoe. <i>Phytochemistry</i> , 2021 , 190, 112843	4	3
501	Plants upcycle gene functions to suit their roots. <i>Trends in Plant Science</i> , 2021 , 26, 996-998	13.1	
500	The knowns and unknowns of intracellular partitioning of carbon and nitrogen, with focus on the organic acid-mediated interplay between mitochondrion and chloroplast. <i>Journal of Plant Physiology</i> , 2021 , 266, 153521	3.6	2
499	Acclimation in plants - the Green Hub consortium. <i>Plant Journal</i> , 2021 , 106, 23-40	6.9	8
498	Ancestral sequence reconstruction - An underused approach to understand the evolution of gene function in plants?. <i>Computational and Structural Biotechnology Journal</i> , 2021 , 19, 1579-1594	6.8	4
497	Comparative Molecular and Metabolic Profiling of Two Contrasting Wheat Cultivars under Drought Stress <i>International Journal of Molecular Sciences</i> , 2021 , 22,	6.3	1
496	Jujube metabolome selection determined the edible properties acquired during domestication. <i>Plant Journal</i> , 2021 ,	6.9	5
495	Selection of a subspecies-specific diterpene gene cluster implicated in rice disease resistance. <i>Nature Plants</i> , 2020 , 6, 1447-1454	11.5	25
494	Co-Regulation of Clustered and Neo-Functionalized Genes in Plant-Specialized Metabolism. <i>Plants</i> , 2020 , 9,	4.5	4
493	An improved extraction method enables the comprehensive analysis of lipids, proteins, metabolites and phytohormones from a single sample of leaf tissue under water-deficit stress. <i>Plant Journal</i> , 2020 , 103, 1614-1632	6.9	16
492	An Abundance and Interaction Encyclopedia of Plant Protein Function. <i>Trends in Plant Science</i> , 2020 , 25, 627-630	13.1	4
491	A NAC transcription factor and its interaction protein hinder abscisic acid biosynthesis by synergistically repressing NCED5 in Citrus reticulata. <i>Journal of Experimental Botany</i> , 2020 , 71, 3613-36	25	17
490	The Kernel Size-Related Quantitative Trait Locus Encodes a Pentatricopeptide Repeat Protein That Aaffects Photosynthesis and Grain Filling. <i>Plant Physiology</i> , 2020 , 183, 1696-1709	6.6	10

489	Phytochrome-Dependent Temperature Perception Modulates Isoprenoid Metabolism. <i>Plant Physiology</i> , 2020 , 183, 869-882	6.6	9
488	Network-based strategies in metabolomics data analysis and interpretation: from molecular networking to biological interpretation. <i>Expert Review of Proteomics</i> , 2020 , 17, 243-255	4.2	30
487	PlantaSyst: Teaming up for Systems Biology and Biotechnology. <i>Trends in Plant Science</i> , 2020 , 25, 621-6	2 43.1	
486	Quantitative trait loci analysis of seed-specialized metabolites reveals seed-specific flavonols and differential regulation of glycoalkaloid content in tomato. <i>Plant Journal</i> , 2020 , 103, 2007-2024	6.9	14
485	The -dependent signalling pathway coordinates plastid biogenesis with the synthesis of anthocyanins. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2020 , 375, 20190403	5.8	8
484	The Cassava Source-Sink project: opportunities and challenges for crop improvement by metabolic engineering. <i>Plant Journal</i> , 2020 , 103, 1655-1665	6.9	11
483	MicroTom Metabolic Network: Rewiring Tomato Metabolic Regulatory Network throughout the Growth Cycle. <i>Molecular Plant</i> , 2020 , 13, 1203-1218	14.4	27
482	Single-Cell Genomics and Epigenomics: Technologies and Applications in Plants. <i>Trends in Plant Science</i> , 2020 , 25, 1030-1040	13.1	19
481	The evolution of metabolism: How to test evolutionary hypotheses at the genomic level. <i>Computational and Structural Biotechnology Journal</i> , 2020 , 18, 482-500	6.8	17
480	Manipulation of ZDS in tomato exposes carotenoid- and ABA-specific effects on fruit development and ripening. <i>Plant Biotechnology Journal</i> , 2020 , 18, 2210-2224	11.6	21
479	Systems-Based Approaches to Unravel Networks and Individual Elements Involved in Apple Superficial Scald. <i>Frontiers in Plant Science</i> , 2020 , 11, 8	6.2	8
478	Type I H+-pyrophosphatase regulates the vacuolar storage of sucrose in citrus fruit. <i>Journal of Experimental Botany</i> , 2020 , 71, 5935-5947	7	2
477	A push, and a pull, to enhance nitrogen use efficiency in rice. <i>Plant Journal</i> , 2020 , 103, 5-6	6.9	O
476	Towards model-driven characterization and manipulation of plant lipid metabolism. <i>Progress in Lipid Research</i> , 2020 , 80, 101051	14.3	9
475	A Highly Efficient -Mediated Method for Transient Gene Expression and Functional Studies in Multiple Plant Species. <i>Plant Communications</i> , 2020 , 1, 100028	9	33
474	Low-temperature tolerance of the Antarctic species Deschampsia antarctica: A complex metabolic response associated with nutrient remobilization. <i>Plant, Cell and Environment</i> , 2020 , 43, 1376-1393	8.4	11
473	Synchronization of developmental, molecular and metabolic aspects of source-sink interactions. <i>Nature Plants</i> , 2020 , 6, 55-66	11.5	52
472	Downregulation of a Mitochondrial NAD+ Transporter (NDT2) Alters Seed Production and Germination in Arabidopsis. <i>Plant and Cell Physiology</i> , 2020 , 61, 897-908	4.9	11

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471	Synthetic analogues of 2-oxo acids discriminate metabolic contribution of the 2-oxoglutarate and 2-oxoadipate dehydrogenases in mammalian cells and tissues. <i>Scientific Reports</i> , 2020 , 10, 1886	4.9	10
470	Metabolomics analysis and metabolite-agronomic trait associations using kernels of wheat (Triticum aestivum) recombinant inbred lines. <i>Plant Journal</i> , 2020 , 103, 279-292	6.9	30
469	Targeting Key Genes to Tailor Old and New Crops for a Greener Agriculture. <i>Molecular Plant</i> , 2020 , 13, 354-356	14.4	4
468	Metabolite-based genome-wide association study enables dissection of the flavonoid decoration pathway of wheat kernels. <i>Plant Biotechnology Journal</i> , 2020 , 18, 1722-1735	11.6	40
467	Metabolic profiles of six African cultivars of cassava (Manihot esculenta Crantz) highlight bottlenecks of root yield. <i>Plant Journal</i> , 2020 , 102, 1202-1219	6.9	17
466	How do vascular plants perform photosynthesis in extreme environments? An integrative ecophysiological and biochemical story. <i>Plant Journal</i> , 2020 , 101, 979-1000	6.9	15
465	Expression Atlas of Provides Insights into the Evolution of Vasculature, Secondary Metabolism, and Roots. <i>Plant Cell</i> , 2020 , 32, 853-870	11.6	17
464	Flux balance analysis of metabolism during growth by osmotic cell expansion and its application to tomato fruits. <i>Plant Journal</i> , 2020 , 103, 68-82	6.9	9
463	Metabolomics in the Context of Plant Natural Products Research: From Sample Preparation to Metabolite Analysis. <i>Metabolites</i> , 2020 , 10,	5.6	58
462	Plant Mitochondrial Carriers: Molecular Gatekeepers That Help to Regulate Plant Central Carbon Metabolism. <i>Plants</i> , 2020 , 9,	4.5	11
461	A Biostimulant Obtained from the Seaweed Protects from Severe Oxidative Stress. <i>International Journal of Molecular Sciences</i> , 2020 , 21,	6.3	20
460	Eating Away at ROS to Regulate Stomatal Opening. <i>Trends in Plant Science</i> , 2020 , 25, 220-223	13.1	18
459	Dissection of flag leaf metabolic shifts and their relationship with those occurring simultaneously in developing seed by application of non-targeted metabolomics. <i>PLoS ONE</i> , 2020 , 15, e0227577	3.7	5
458	Ascorbate and Thiamin: Metabolic Modulators in Plant Acclimation Responses. <i>Plants</i> , 2020 , 9,	4.5	8
457	Lipidomic and transcriptomic analysis reveals reallocation of carbon flux from cuticular wax into plastid membrane lipids in a glossy "Newhall" navel orange mutant. <i>Horticulture Research</i> , 2020 , 7, 41	7.7	6
456	Nano and Micro Unmanned Aerial Vehicles (UAVs): A New Grand Challenge for Precision Agriculture?. <i>Current Protocols in Plant Biology</i> , 2020 , 5, e20103	2.8	6
455	The style and substance of plant flavonoid decoration; towards defining both structure and function. <i>Phytochemistry</i> , 2020 , 174, 112347	4	45
454	Metabolome Profiling Supports the Key Role of the Spike in Wheat Yield Performance. <i>Cells</i> , 2020 , 9,	7.9	9

453	High-Throughput CRISPR/Cas9 Mutagenesis Streamlines Trait Gene Identification in Maize. <i>Plant Cell</i> , 2020 , 32, 1397-1413	11.6	68
452	Analysis of Tomato Post-Harvest Properties: Fruit Color, Shelf Life, and Fungal Susceptibility. <i>Current Protocols in Plant Biology</i> , 2020 , 5, e20108	2.8	5
45 ¹	Outstanding questions in flower metabolism. <i>Plant Journal</i> , 2020 , 103, 1275-1288	6.9	6
450	Resolving the metabolon: is the proof in the metabolite?. <i>EMBO Reports</i> , 2020 , 21, e50774	6.5	3
449	The Penium margaritaceum Genome: Hallmarks of the Origins of Land Plants. Cell, 2020, 181, 1097-111	15612	62
448	Conservation and diversification of flavonoid metabolism in the plant kingdom. <i>Current Opinion in Plant Biology</i> , 2020 , 55, 100-108	9.9	37
447	Manipulation of Etarotene levels in tomato fruits results in increased ABA content and extended shelf life. <i>Plant Biotechnology Journal</i> , 2020 , 18, 1185-1199	11.6	36
446	Construction and applications of a B vitamin genetic resource for investigation of vitamin-dependent metabolism in maize. <i>Plant Journal</i> , 2020 , 101, 442-454	6.9	5
445	Genome-wide Dissection of Co-selected UV-B Responsive Pathways in the UV-B Adaptation of Qingke. <i>Molecular Plant</i> , 2020 , 13, 112-127	14.4	42
444	Metabolomics should be deployed in the identification and characterization of gene-edited crops. <i>Plant Journal</i> , 2020 , 102, 897-902	6.9	24
443	The Acetate Pathway Supports Flavonoid and Lipid Biosynthesis in Arabidopsis. <i>Plant Physiology</i> , 2020 , 182, 857-869	6.6	19
442	Wasteful, essential, evolutionary stepping stone? The multiple personalities of the photorespiratory pathway. <i>Plant Journal</i> , 2020 , 102, 666-677	6.9	20
441	Assessing durum wheat ear and leaf metabolomes in the field through hyperspectral data. <i>Plant Journal</i> , 2020 , 102, 615-630	6.9	18
440	A genetically validated approach for detecting inorganic polyphosphates in plants. <i>Plant Journal</i> , 2020 , 102, 507-516	6.9	7
439	The Past, Present, and Future of Maize Improvement: Domestication, Genomics, and Functional Genomic Routes toward Crop Enhancement. <i>Plant Communications</i> , 2020 , 1, 100010	9	34
438	Full-Length Transcript-Based Proteogenomics of Rice Improves Its Genome and Proteome Annotation. <i>Plant Physiology</i> , 2020 , 182, 1510-1526	6.6	30
437	Changes in intracellular NAD status affect stomatal development in an abscisic acid-dependent manner. <i>Plant Journal</i> , 2020 , 104, 1149-1168	6.9	7
436	Mobile Transposable Elements Shape Plant Genome Diversity. <i>Trends in Plant Science</i> , 2020 , 25, 1062-10) 64 .1	3

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435	Model-assisted identification of metabolic engineering strategies for Jatropha curcas lipid pathways. <i>Plant Journal</i> , 2020 , 104, 76-95	6.9	7
434	How do wheat plants cope with Pyricularia oryzae infection? A physiological and metabolic approach. <i>Planta</i> , 2020 , 252, 24	4.7	1
433	Born to revive: molecular and physiological mechanisms of double tolerance in a paleotropical and resurrection plant. <i>New Phytologist</i> , 2020 , 226, 741-759	9.8	17
432	Plant Single-Cell Metabolomics-Challenges and Perspectives. <i>International Journal of Molecular Sciences</i> , 2020 , 21,	6.3	9
431	Role of Raf-like kinases in SnRK2 activation and osmotic stress response in plants. <i>Nature Communications</i> , 2020 , 11, 6184	17.4	18
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408 407		1.4	1
	Spectrometry Metabolomics. <i>Methods in Molecular Biology</i> , 2020 , 2104, 209-225 Dissection of flag leaf metabolic shifts and their relationship with those occurring simultaneously in	1.4	1
407	Spectrometry Metabolomics. <i>Methods in Molecular Biology</i> , 2020 , 2104, 209-225 Dissection of flag leaf metabolic shifts and their relationship with those occurring simultaneously in developing seed by application of non-targeted metabolomics 2020 , 15, e0227577 Dissection of flag leaf metabolic shifts and their relationship with those occurring simultaneously in	1.4	1
4 ⁰ 7	Dissection of flag leaf metabolic shifts and their relationship with those occurring simultaneously in developing seed by application of non-targeted metabolomics 2020, 15, e0227577 Dissection of flag leaf metabolic shifts and their relationship with those occurring simultaneously in developing seed by application of non-targeted metabolomics 2020, 15, e0227577 Dissection of flag leaf metabolic shifts and their relationship with those occurring simultaneously in Dissection of flag leaf metabolic shifts and their relationship with those occurring simultaneously in	1.4	1
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407 406 405 404	Dissection of flag leaf metabolic shifts and their relationship with those occurring simultaneously in developing seed by application of non-targeted metabolomics 2020, 15, e0227577 Dissection of flag leaf metabolic shifts and their relationship with those occurring simultaneously in developing seed by application of non-targeted metabolomics 2020, 15, e0227577 Dissection of flag leaf metabolic shifts and their relationship with those occurring simultaneously in developing seed by application of non-targeted metabolomics 2020, 15, e0227577 Dissection of flag leaf metabolic shifts and their relationship with those occurring simultaneously in developing seed by application of non-targeted metabolomics 2020, 15, e0227577 Redox-Regulation of Photorespiration through Mitochondrial Thioredoxin o1. <i>Plant Physiology</i> ,		
407 406 405 404 403	Dissection of flag leaf metabolic shifts and their relationship with those occurring simultaneously in developing seed by application of non-targeted metabolomics 2020, 15, e0227577 Dissection of flag leaf metabolic shifts and their relationship with those occurring simultaneously in developing seed by application of non-targeted metabolomics 2020, 15, e0227577 Dissection of flag leaf metabolic shifts and their relationship with those occurring simultaneously in developing seed by application of non-targeted metabolomics 2020, 15, e0227577 Dissection of flag leaf metabolic shifts and their relationship with those occurring simultaneously in developing seed by application of non-targeted metabolomics 2020, 15, e0227577 Redox-Regulation of Photorespiration through Mitochondrial Thioredoxin o1. <i>Plant Physiology</i> , 2019, 181, 442-457 Pan-Genomic Illumination of Tomato Identifies Novel Gene-Trait Interactions. <i>Trends in Plant</i>	6.6	25

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300 299		7	157
	of Experimental Botany, 2017 , 68, 4013-4028		
299	of Experimental Botany, 2017, 68, 4013-4028 Canalization of Tomato Fruit Metabolism. Plant Cell, 2017, 29, 2753-2765 Floral Metabolism of Sugars and Amino Acids: Implications for Pollinators/Preferences and Seed	11.6	34
299 298	of Experimental Botany, 2017, 68, 4013-4028 Canalization of Tomato Fruit Metabolism. Plant Cell, 2017, 29, 2753-2765 Floral Metabolism of Sugars and Amino Acids: Implications for Pollinators/Preferences and Seed and Fruit Set. Plant Physiology, 2017, 175, 1510-1524	11.6	34 50
299 298 297	Canalization of Tomato Fruit Metabolism. <i>Plant Cell</i> , 2017 , 29, 2753-2765 Floral Metabolism of Sugars and Amino Acids: Implications for PollinatorsNPreferences and Seed and Fruit Set. <i>Plant Physiology</i> , 2017 , 175, 1510-1524 Metabolism within the specialized guard cells of plants. <i>New Phytologist</i> , 2017 , 216, 1018-1033 The Photorespiratory Metabolite 2-Phosphoglycolate Regulates Photosynthesis and Starch	11.6 6.6 9.8 11.6	34 50 42
299 298 297 296	Canalization of Tomato Fruit Metabolism. <i>Plant Cell</i> , 2017 , 29, 2753-2765 Floral Metabolism of Sugars and Amino Acids: Implications for PollinatorsNPreferences and Seed and Fruit Set. <i>Plant Physiology</i> , 2017 , 175, 1510-1524 Metabolism within the specialized guard cells of plants. <i>New Phytologist</i> , 2017 , 216, 1018-1033 The Photorespiratory Metabolite 2-Phosphoglycolate Regulates Photosynthesis and Starch Accumulation in Arabidopsis. <i>Plant Cell</i> , 2017 , 29, 2537-2551	11.6 6.6 9.8 11.6	34504271
299 298 297 296	Canalization of Tomato Fruit Metabolism. <i>Plant Cell</i> , 2017 , 29, 2753-2765 Floral Metabolism of Sugars and Amino Acids: Implications for PollinatorsNPreferences and Seed and Fruit Set. <i>Plant Physiology</i> , 2017 , 175, 1510-1524 Metabolism within the specialized guard cells of plants. <i>New Phytologist</i> , 2017 , 216, 1018-1033 The Photorespiratory Metabolite 2-Phosphoglycolate Regulates Photosynthesis and Starch Accumulation in Arabidopsis. <i>Plant Cell</i> , 2017 , 29, 2537-2551 De Novo Assembly of a New Accession Using Nanopore Sequencing. <i>Plant Cell</i> , 2017 , 29, 2336-2348	11.6 6.6 9.8 11.6	34 50 42 71 138

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	Metabolic variation between japonica and indica rice cultivars as revealed by non-targeted		
209	Metabolic variation between japonica and indica rice cultivars as revealed by non-targeted metabolomics. <i>Scientific Reports</i> , 2014 , 4, 5067 Silencing of the tomato sugar partitioning affecting protein (SPA) modifies sink strength through a	4.9	109
209	Metabolic variation between japonica and indica rice cultivars as revealed by non-targeted metabolomics. <i>Scientific Reports</i> , 2014 , 4, 5067 Silencing of the tomato sugar partitioning affecting protein (SPA) modifies sink strength through a shift in leaf sugar metabolism. <i>Plant Journal</i> , 2014 , 77, 676-87 Genome-enabled plant metabolomics. <i>Journal of Chromatography B: Analytical Technologies in the</i>	4.9 6.9	109
209 208	Metabolic variation between japonica and indica rice cultivars as revealed by non-targeted metabolomics. <i>Scientific Reports</i> , 2014 , 4, 5067 Silencing of the tomato sugar partitioning affecting protein (SPA) modifies sink strength through a shift in leaf sugar metabolism. <i>Plant Journal</i> , 2014 , 77, 676-87 Genome-enabled plant metabolomics. <i>Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences</i> , 2014 , 966, 7-20 Conserved changes in the dynamics of metabolic processes during fruit development and ripening	4.9 6.9 3.2	109 15 34
209208207206	Metabolic variation between japonica and indica rice cultivars as revealed by non-targeted metabolomics. <i>Scientific Reports</i> , 2014 , 4, 5067 Silencing of the tomato sugar partitioning affecting protein (SPA) modifies sink strength through a shift in leaf sugar metabolism. <i>Plant Journal</i> , 2014 , 77, 676-87 Genome-enabled plant metabolomics. <i>Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences</i> , 2014 , 966, 7-20 Conserved changes in the dynamics of metabolic processes during fruit development and ripening across species. <i>Plant Physiology</i> , 2014 , 164, 55-68 Mercator: a fast and simple web server for genome scale functional annotation of plant sequence	4.9 6.9 3.2 6.6	109 15 34 42
209208207206205	Metabolic variation between japonica and indica rice cultivars as revealed by non-targeted metabolomics. <i>Scientific Reports</i> , 2014 , 4, 5067 Silencing of the tomato sugar partitioning affecting protein (SPA) modifies sink strength through a shift in leaf sugar metabolism. <i>Plant Journal</i> , 2014 , 77, 676-87 Genome-enabled plant metabolomics. <i>Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences</i> , 2014 , 966, 7-20 Conserved changes in the dynamics of metabolic processes during fruit development and ripening across species. <i>Plant Physiology</i> , 2014 , 164, 55-68 Mercator: a fast and simple web server for genome scale functional annotation of plant sequence data. <i>Plant, Cell and Environment</i> , 2014 , 37, 1250-8 Comparative analyses of Cland Cland Cland comparative and rice. <i>Nature</i>	4.9 6.9 3.2 6.6 8.4	109 15 34 42 373

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120	Genetics, Genomics and Metabolomics 2011 , 219-259 Metabolic priming by a secreted fungal effector. <i>Nature</i> , 2011 , 478, 395-8	50.4	5 371
		50.4	371
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119 118 117	Metabolic priming by a secreted fungal effector. <i>Nature</i> , 2011 , 478, 395-8 Recommendations for reporting metabolite data. <i>Plant Cell</i> , 2011 , 23, 2477-82 Genetic dissection of vitamin E biosynthesis in tomato. <i>Journal of Experimental Botany</i> , 2011 , 62, 3781-	11.6 98	371 238 58
119 118 117	Metabolic priming by a secreted fungal effector. <i>Nature</i> , 2011 , 478, 395-8 Recommendations for reporting metabolite data. <i>Plant Cell</i> , 2011 , 23, 2477-82 Genetic dissection of vitamin E biosynthesis in tomato. <i>Journal of Experimental Botany</i> , 2011 , 62, 3781- Fumarate: Multiple functions of a simple metabolite. <i>Phytochemistry</i> , 2011 , 72, 838-43 Transcriptome and metabolite profiling show that APETALA2a is a major regulator of tomato fruit	11.6 98 ³	3712385855
119 118 117 116	Metabolic priming by a secreted fungal effector. <i>Nature</i> , 2011 , 478, 395-8 Recommendations for reporting metabolite data. <i>Plant Cell</i> , 2011 , 23, 2477-82 Genetic dissection of vitamin E biosynthesis in tomato. <i>Journal of Experimental Botany</i> , 2011 , 62, 3781- Fumarate: Multiple functions of a simple metabolite. <i>Phytochemistry</i> , 2011 , 72, 838-43 Transcriptome and metabolite profiling show that APETALA2a is a major regulator of tomato fruit ripening. <i>Plant Cell</i> , 2011 , 23, 923-41 Metabolic profiling during peach fruit development and ripening reveals the metabolic networks	11.6 98 4 11.6	3712385855269

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5	Pod indehiscence in common bean is associated to the fine regulation of PvMYB26 and a non-functional abscission layer		4
4	The Genome of the Charophyte AlgaPenium margaritaceumBears Footprints of the Evolutionary Origins of Land Plants		3

3 Portrait of a genus: the genetic diversity of Zea

2	Regulation of Plant Primary Metabolism [How Results From Novel Technologies Are Extending Our Understanding From Classical Targeted Approaches. <i>Critical Reviews in Plant Sciences</i> ,1-20	5.6	О
1	Genetic architecture of seed glycerolipids in Asian cultivated rice. Plant, Cell and Environment,	8.4	3

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