Richard Wilson

List of Publications by Year in descending order

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221 papers 10,168 citations

53 h-index 89 g-index

230 all docs

230 docs citations

times ranked

230

9417 citing authors

#	Article	IF	CITATIONS
1	Cadmium-induced changes in composition and co-metabolism of glycerolipids species in wheat root: Glycerolipidomic and transcriptomic approach. Journal of Hazardous Materials, 2022, 423, 127115.	12.4	8
2	Reduced apoplastic barriers in tissues of shoot-proximal rhizomes of <i>Oryza coarctata</i> are associated with Na+ sequestration. Journal of Experimental Botany, 2022, 73, 998-1015.	4.8	2
3	Nucleotide-binding leucine-rich repeat proteins: a missing link in controlling cell fate and plant adaptation to hostile environment?. Journal of Experimental Botany, 2022, 73, 631-635.	4.8	1
4	Cation transporters in cell fate determination and plant adaptive responses to a low-oxygen environment. Journal of Experimental Botany, 2022, 73, 636-645.	4.8	7
5	Genome-wide association study reveals a genomic region on 5AL for salinity tolerance in wheat. Theoretical and Applied Genetics, 2022, 135, 709-721.	3.6	10
6	Rewilding staple crops for the lost halophytism: Toward sustainability and profitability of agricultural production systems. Molecular Plant, 2022, 15, 45-64.	8.3	23
7	Shotgun Proteomics as a Powerful Tool for the Study of the Proteomes of Plants, Their Pathogens, and Plant–Pathogen Interactions. Proteomes, 2022, 10, 5.	3.5	18
8	Proto Kranz-like leaf traits and cellular ionic regulation are associated with salinity tolerance in a halophytic wild rice. Stress Biology, 2022, 2, 1.	3.1	4
9	Proteomic investigation of brain, liver and intestine in high feed intake and low feed intake Chinook salmon (Oncorhynchus tshawytscha). Aquaculture, 2022, 551, 737915.	3.5	9
10	Impacts of barley root cortical aerenchyma on growth, physiology, yield components, and grain quality under field waterlogging conditions. Field Crops Research, 2022, 279, 108461.	5.1	9
11	The role of NADPH oxidases in regulating leaf gas exchange and ion homeostasis in Arabidopsis plants under cadmium stress. Journal of Hazardous Materials, 2022, 429, 128217.	12.4	11
12	Application of omics technologies in single-type guard cell studies for understanding the mechanistic basis of plant adaptation to saline conditions. Advances in Botanical Research, 2022, , 249-270.	1.1	2
13	Comparative Analysis of Root Na+ Relation under Salinity between OryzaÂsativa and Oryza coarctata. Plants, 2022, 11, 656.	3.5	7
14	Rethinking Rehabilitation of Salt-Affected Land: New Perspectives from Australian Experience. Earth, 2022, 3, 245-258.	2.2	3
15	Using Portable X-ray Fluorescence Spectroscopy for Inexpensive and Quick Determination of Micronutrients in Barley Shoots. Communications in Soil Science and Plant Analysis, 2022, 53, 1379-1384.	1.4	4
16	Evolutionary Significance of NHX Family and NHX1 in Salinity Stress Adaptation in the Genus Oryza. International Journal of Molecular Sciences, 2022, 23, 2092.	4.1	19
17	Plant responses to heterogeneous salinity: agronomic relevance and research priorities. Annals of Botany, 2022, 129, 499-518.	2.9	13
18	Cathelicidin-3 Associated With Serum Extracellular Vesicles Enables Early Diagnosis of a Transmissible Cancer. Frontiers in Immunology, 2022, 13, 858423.	4.8	3

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19	Genome-Wide Association Study Reveals Marker Trait Associations (MTA) for Waterlogging-Triggered Adventitious Roots and Aerenchyma Formation in Barley. International Journal of Molecular Sciences, 2022, 23, 3341.	4.1	9
20	Multidimensional screening and evaluation of morphoâ€physiological indices for salinity stress tolerance in wheat. Journal of Agronomy and Crop Science, 2022, 208, 454-471.	3. 5	11
21	pH-Dependent mitigation of aluminum toxicity in pea (Pisum sativum) roots by boron. Plant Science, 2022, 318, 111208.	3.6	7
22	Challenges of an Emerging Disease: The Evolving Approach to Diagnosing Devil Facial Tumour Disease. Pathogens, 2022, 11, 27.	2.8	1
23	Large-Scale Protein and Phosphoprotein Profiling to Explore Potato Resistance Mechanisms to Spongospora subterranea Infection. Frontiers in Plant Science, 2022, 13, 872901.	3.6	3
24	A novel R3H protein, OsDIP1, confers ABA-mediated adaptation to drought and salinity stress in rice. Plant and Soil, 2022, 477, 501-519.	3.7	1
25	Local and systemic responses conferring acclimation of <i>Brassica napus</i> roots to low phosphorus conditions. Journal of Experimental Botany, 2022, 73, 4753-4777.	4.8	9
26	Tissue-Specific Responses of Cereals to Two Fusarium Diseases and Effects of Plant Height and Drought Stress on Their Susceptibility. Agronomy, 2022, 12, 1108.	3.0	3
27	Stalk cell polar ion transport provide for bladderâ€based salinity tolerance in <i>Chenopodium quinoa</i> . New Phytologist, 2022, 235, 1822-1835.	7.3	8
28	Evaluation of salt tolerance of oat cultivars and the mechanism of adaptation to salinity. Journal of Plant Physiology, 2022, 273, 153708.	3.5	15
29	Protein metabolism in the liver and white muscle is associated with feed efficiency in Chinook salmon (Oncorhynchus tshawytscha) reared in seawater: Evidence from proteomic analysis. Comparative Biochemistry and Physiology Part D: Genomics and Proteomics, 2022, 42, 100994.	1.0	2
30	Melatonin as a regulator of plant ionic homeostasis: implications for abiotic stress tolerance. Journal of Experimental Botany, 2022, 73, 5886-5902.	4.8	26
31	Multi-omics reveals mechanisms of resistance to potato root infection by Spongospora subterranea. Scientific Reports, 2022, 12, .	3.3	6
32	Non-stomatal limitation of photosynthesis by soil salinity. Critical Reviews in Environmental Science and Technology, 2021, 51, 791-825.	12.8	129
33	Evolution of rapid blueâ€light response linked to explosive diversification of ferns in angiosperm forests. New Phytologist, 2021, 230, 1201-1213.	7.3	33
34	A comparative analysis of stomatal traits and photosynthetic responses in closely related halophytic and glycophytic species under saline conditions. Environmental and Experimental Botany, 2021, 181, 104300.	4.2	36
35	Salinity Effects on Guard Cell Proteome in Chenopodium quinoa. International Journal of Molecular Sciences, 2021, 22, 428.	4.1	20
36	Antioxidant Enzymatic Activity and Osmotic Adjustment as Components of the Drought Tolerance Mechanism in Carex duriuscula. Plants, 2021, 10, 436.	3 . 5	25

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37	Sodium sequestration confers salinity tolerance in an ancestral wild rice. Physiologia Plantarum, 2021, 172, 1594-1608.	5.2	22
38	Biochemical and biophysical pH clamp controlling Net H ⁺ efflux across the plasma membrane of plant cells. New Phytologist, 2021, 230, 408-415.	7.3	25
39	Optimizing hydroponic culture media and NO3â [^] /NH4+ ratio for improving essential oil compositions of purple coneflower (Echinacea purpurea L.). Scientific Reports, 2021, 11, 8009.	3.3	8
40	Quantitative proteomics provides an insight into germinationâ€related proteins in the obligate biotrophic plant pathogen <scp><i>Spongospora subterranea</i></scp> . Environmental Microbiology Reports, 2021, 13, 521-532.	2.4	9
41	Understanding the mechanistic basis of adaptation of perennial <i>Sarcocornia quinqueflora</i> species to soil salinity. Physiologia Plantarum, 2021, 172, 1997-2010.	5.2	18
42	Hypoxia-induced increase in GABA content is essential for restoration of membrane potential and preventing ROS-induced disturbance to ion homeostasis. Plant Communications, 2021, 2, 100188.	7.7	47
43	Early responses to salt stress in quinoa genotypes with opposite behavior. Physiologia Plantarum, 2021, 173, 1392-1420.	5.2	10
44	Improving Performance of Salt-Grown Crops by Exogenous Application of Plant Growth Regulators. Biomolecules, 2021, 11, 788.	4.0	46
45	Avenues for biofortification of zinc in barley for human and animal health: a meta-analysis. Plant and Soil, 2021, 466, 101-119.	3.7	5
46	Rewilding crops for climate resilience: economic analysis and <i>de novo</i> domestication strategies. Journal of Experimental Botany, 2021, 72, 6123-6139.	4.8	52
47	Revealing the Role of the Calcineurin B-Like Protein-Interacting Protein Kinase 9 (CIPK9) in Rice Adaptive Responses to Salinity, Osmotic Stress, and K+ Deficiency. Plants, 2021, 10, 1513.	3.5	9
48	Effects of TDP-43 overexpression on neuron proteome and morphology in vitro. Molecular and Cellular Neurosciences, 2021, 114, 103627.	2.2	1
49	Increasing medicinal and phytochemical compounds of coneflower (Echinacea purpurea L.) as affected by NO3â°/NH4+ ratio and perlite particle size in hydroponics. Scientific Reports, 2021, 11, 15202.	3.3	7
50	AFB1 controls rapid auxin signalling through membrane depolarization in Arabidopsis thaliana root. Nature Plants, 2021, 7, 1229-1238.	9.3	59
51	In Planta Transcriptome and Proteome Profiles of Spongospora subterranea in Resistant and Susceptible Host Environments Illuminates Regulatory Principles Underlying Host–Pathogen Interaction. Biology, 2021, 10, 840.	2.8	8
52	Early signalling processes in roots play a crucial role in the differential salt tolerance in contrasting Chenopodium quinoa accessions. Journal of Experimental Botany, 2021, , .	4.8	4
53	Tissue-specificity of ROS-induced K+ and Ca2+ fluxes in succulent stems of the perennial halophyte Sarcocornia quinqueflora in the context of salinity stress tolerance. Plant Physiology and Biochemistry, 2021, 166, 1022-1031.	5.8	7
54	Understanding a Mechanistic Basis of ABA Involvement in Plant Adaptation to Soil Flooding: The Current Standing. Plants, 2021, 10, 1982.	3.5	16

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55	Properties of an acid-tolerant, persistent Cheddar cheese isolate, <i>Lacticaseibacillus paracasei</i> GCRL163. Journal of Industrial Microbiology and Biotechnology, 2021, 48, .	3.0	O
56	Phosphoinositides: Emerging players in plant salinity stress tolerance. Molecular Plant, 2021, 14, 1973-1975.	8.3	2
57	Mitochondrial respiratory chain function promotes extracellular matrix integrity in cartilage. Journal of Biological Chemistry, 2021, 297, 101224.	3.4	16
58	Proteomic investigation of liver and white muscle in efficient and inefficient Chinook salmon (Oncorhynchus tshawytscha): Fatty acid metabolism and protein turnover drive feed efficiency. Aquaculture, 2021, 542, 736855.	3.5	24
59	Ionomics analysis provides new insights into the co-enrichment of cadmium and zinc in wheat grains. Ecotoxicology and Environmental Safety, 2021, 223, 112623.	6.0	16
60	Arsenic transport and interaction with plant metabolism: Clues for improving agricultural productivity and food safety. Environmental Pollution, 2021, 290, 117987.	7.5	54
61	Protein levels, air pollution and vitamin D deficiency: links with allergy. ERJ Open Research, 2021, 7, 00237-2021.	2.6	0
62	Extracellular vesicle proteomes of two transmissible cancers of Tasmanian devils reveal tenascin-C as a serum-based differential diagnostic biomarker. Cellular and Molecular Life Sciences, 2021, 78, 7537-7555.	5.4	6
63	Cell surface and intracellular auxin signalling for H+ fluxes in root growth. Nature, 2021, 599, 273-277.	27.8	128
64	Tissue tolerance mechanisms conferring salinity tolerance in a halophytic perennial species <i>Nitraria sibirica </i> Pall Tree Physiology, 2021, 41, 1264-1277.	3.1	22
65	Effects of Potassium Availability on Growth and Development of Barley Cultivars. Agronomy, 2021, 11, 2269.	3.0	6
66	To exclude or to accumulate? Revealing the role of the sodium HKT1;5 transporter in plant adaptive responses to varying soil salinity. Plant Physiology and Biochemistry, 2021, 169, 333-342.	5.8	20
67	Doing â€~business as usual' comes with a cost: evaluating energy cost of maintaining plant intracellular K ⁺ homeostasis under saline conditions. New Phytologist, 2020, 225, 1097-1104.	7.3	140
68	Energy costs of salt tolerance in crop plants. New Phytologist, 2020, 225, 1072-1090.	7.3	284
69	The energy cost of the tonoplast futile sodium leak. New Phytologist, 2020, 225, 1105-1110.	7.3	86
70	Biochemical pH clamp: the forgotten resource in membrane bioenergetics. New Phytologist, 2020, 225, 37-47.	7.3	33
71	Two of a kind: transmissible Schwann cell cancers in the endangered Tasmanian devil (Sarcophilus) Tj ETQq $1\ 1\ 0$.784314 r 5.4	gBT /Overloc
72	Identification of Key Pro-Survival Proteins in Isolated Colonic Goblet Cells of Winnie, a Murine Model of Spontaneous Colitis. Inflammatory Bowel Diseases, 2020, 26, 80-92.	1.9	5

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73	Comparing Kinetics of Xylem Ion Loading and Its Regulation in Halophytes and Glycophytes. Plant and Cell Physiology, 2020, 61, 403-415.	3.1	22
74	Stomatal traits as a determinant of superior salinity tolerance in wild barley. Journal of Plant Physiology, 2020, 245, 153108.	3.5	41
75	A hemeâ€binding protein produced by <i>Haemophilus haemolyticus</i> inhibits nonâ€typeable <i>Haemophilus influenzae</i> . Molecular Microbiology, 2020, 113, 381-398.	2.5	18
76	Phylogenetic Diversity and Physiological Roles of Plant Monovalent Cation/H+ Antiporters. Frontiers in Plant Science, 2020, 11, 573564.	3.6	45
77	NADPH oxidases and the evolution of plant salinity tolerance. Plant, Cell and Environment, 2020, 43, 2957-2968.	5.7	49
78	What makes a plant science manuscript successful for publication?. Functional Plant Biology, 2020, 47, 1138.	2.1	3
79	Lipid kinases PIP5K7 and PIP5K9 are required for polyamineâ€triggered K ⁺ efflux in Arabidopsis roots. Plant Journal, 2020, 104, 416-432.	5.7	28
80	The proteomic response is linked to regional lung volumes in ventilator-induced lung injury. Journal of Applied Physiology, 2020, 129, 837-845.	2.5	6
81	Changes in Expression Level of OsHKT1;5 Alters Activity of Membrane Transporters Involved in K+ and Ca2+ Acquisition and Homeostasis in Salinized Rice Roots. International Journal of Molecular Sciences, 2020, 21, 4882.	4.1	23
82	Optimisation of Sporosori Purification and Protein Extraction Techniques for the Biotrophic Protozoan Plant Pathogen Spongospora subterranea. Molecules, 2020, 25, 3109.	3.8	9
83	A simple apparatus for electrokinetic removal of sodium dodecyl sulfate from protein digests. Journal of Chromatography A, 2020, 1628, 461443.	3.7	1
84	Candidate genes for salinity tolerance in barley revealed by RNA-seq analysis of near-isogenic lines. Plant Growth Regulation, 2020, 92, 571-582.	3.4	14
85	Leaf mesophyll K+ and Clâ^ fluxes and reactive oxygen species production predict rice salt tolerance at reproductive stage in greenhouse and field conditions. Plant Growth Regulation, 2020, 92, 53-64.	3.4	18
86	Understanding the role of root-related traits in salinity tolerance of quinoa accessions with contrasting epidermal bladder cell patterning. Planta, 2020, 251, 103.	3.2	14
87	Homology Modeling Identifies Crucial Amino-Acid Residues That Confer Higher Na+ Transport Capacity of OcHKT1;5 from Oryza coarctata Roxb. Plant and Cell Physiology, 2020, 61, 1321-1334.	3.1	23
88	Calcium-Dependent Hydrogen Peroxide Mediates Hydrogen-Rich Water-Reduced Cadmium Uptake in Plant Roots. Plant Physiology, 2020, 183, 1331-1344.	4.8	34
89	Mechanisms of Plant Responses and Adaptation to Soil Salinity. Innovation(China), 2020, 1, 100017.	9.1	387
90	Developing and validating protocols for mechanical isolation of guard-cell enriched epidermal peels for omics studies. Functional Plant Biology, 2020, 47, 803.	2.1	8

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91	Ablation of the miRNA Cluster 24 Has Profound Effects on Extracellular Matrix Protein Abundance in Cartilage. International Journal of Molecular Sciences, 2020, 21, 4112.	4.1	5
92	Melatonin improves rice salinity stress tolerance by <scp>NADPH</scp> oxidaseâ€dependent control of the plasma membrane K ⁺ transporters and K ⁺ homeostasis. Plant, Cell and Environment, 2020, 43, 2591-2605.	5.7	93
93	Prospects for the accelerated improvement of the resilient crop quinoa. Journal of Experimental Botany, 2020, 71, 5333-5347.	4.8	49
94	Understanding Mechanisms of Salinity Tolerance in Barley by Proteomic and Biochemical Analysis of Near-Isogenic Lines. International Journal of Molecular Sciences, 2020, 21, 1516.	4.1	45
95	Function of NHX-type transporters in improving rice tolerance to aluminum stress and soil acidity. Planta, 2020, 251, 71.	3.2	23
96	Prolonged Heat Stress of Lactobacillus paracasei GCRL163 Improves Binding to Human Colorectal Adenocarcinoma HT-29 Cells and Modulates the Relative Abundance of Secreted and Cell Surface-Located Proteins. Journal of Proteome Research, 2020, 19, 1824-1846.	3.7	7
97	Reducing Cadmium Accumulation in Plants: Structure–Function Relations and Tissue-Specific Operation of Transporters in the Spotlight. Plants, 2020, 9, 223.	3.5	88
98	GORK Channel: A Master Switch of Plant Metabolism?. Trends in Plant Science, 2020, 25, 434-445.	8.8	73
99	Identification of new QTL for salt tolerance from rice variety Pokkali. Journal of Agronomy and Crop Science, 2020, 206, 202-213.	3.5	31
100	Linking phytohormones with growth, transport activity and metabolic responses to cadmium in tomato. Plant Growth Regulation, 2020, 90, 557-569.	3.4	7
101	Distinct Evolutionary Origins of Intron Retention Splicing Events in NHX1 Antiporter Transcripts Relate to Sequence Specific Distinctions in Oryza Species. Frontiers in Plant Science, 2020, 11, 267.	3.6	16
102	Back to the Wild: On a Quest for Donors Toward Salinity Tolerant Rice. Frontiers in Plant Science, 2020, 11, 323.	3.6	54
103	Sugar Beet (Beta vulgaris) Guard Cells Responses to Salinity Stress: A Proteomic Analysis. International Journal of Molecular Sciences, 2020, 21, 2331.	4.1	16
104	Modulation of Ion Transport Across Plant Membranes by Polyamines: Understanding Specific Modes of Action Under Stress. Frontiers in Plant Science, 2020, 11, 616077.	3.6	21
105	Linking sensitivity of photosystem II to UV-B with chloroplast ultrastructure and UV-B absorbing pigments contents in A. thaliana L. phyAphyB double mutants. Plant Growth Regulation, 2020, 91, 13-21.	3.4	13
106	Crop Halophytism: An Environmentally Sustainable Solution for Global Food Security. Trends in Plant Science, 2020, 25, 630-634.	8.8	77
107	The State of the Art in Modeling Waterlogging Impacts on Plants: What Do We Know and What Do We Need to Know. Earth's Future, 2020, 8, e2020EF001801.	6.3	49
108	Hydrogen-rich water promotes elongation of hypocotyls and roots in plants through mediating the level of endogenous gibberellin and auxin. Functional Plant Biology, 2020, 47, 771.	2.1	15

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109	Control of xylem Na ⁺ loading and transport to the shoot in rice and barley as a determinant of differential salinity stress tolerance. Physiologia Plantarum, 2019, 165, 619-631.	5.2	50
110	An RNA-binding protein MUG13.4 interacts with AtAGO2 to modulate salinity tolerance in Arabidopsis. Plant Science, 2019, 288, 110218.	3.6	9
111	Genomic regions on chromosome 5H containing a novel QTL conferring barley yellow dwarf virus-PAV (BYDV-PAV) tolerance in barley. Scientific Reports, 2019, 9, 11298.	3.3	11
112	GABA operates upstream of H+-ATPase and improves salinity tolerance in Arabidopsis by enabling cytosolic K+ retention and Na+ exclusion. Journal of Experimental Botany, 2019, 70, 6349-6361.	4.8	73
113	Tissue-specific respiratory burst oxidase homolog-dependent H2O2 signaling to the plasma membrane H+-ATPase confers potassium uptake and salinity tolerance in Cucurbitaceae. Journal of Experimental Botany, 2019, 70, 5879-5893.	4.8	90
114	Tissue-Specific Regulation of Na+ and K+ Transporters Explains Genotypic Differences in Salinity Stress Tolerance in Rice. Frontiers in Plant Science, 2019, 10, 1361.	3.6	67
115	A large-scale screening of quinoa accessions reveals an important role of epidermal bladder cells and stomatal patterning in salinity tolerance. Environmental and Experimental Botany, 2019, 168, 103885.	4.2	39
116	Extracellular Spermine Triggers a Rapid Intracellular Phosphatidic Acid Response in Arabidopsis, Involving PLDÎ Activation and Stimulating Ion Flux. Frontiers in Plant Science, 2019, 10, 601.	3.6	19
117	Root vacuolar Na ⁺ sequestration but not exclusion from uptake correlates with barley salt tolerance. Plant Journal, 2019, 100, 55-67.	5.7	80
118	Plants Grown in Parafilm-Wrapped Petri Dishes Are Stressed and Possess Altered Gene Expression Profile. Frontiers in Plant Science, 2019, 10, 637.	3.6	14
119	Microhair on the adaxial leaf surface of salt secreting halophytic Oryza coarctata Roxb. show distinct morphotypes: Isolation for molecular and functional analysis. Plant Science, 2019, 285, 248-257.	3.6	16
120	Extracellular silica nanocoat formed by layer-by-layer (LBL) self-assembly confers aluminum resistance in root border cells of pea (Pisum sativum). Journal of Nanobiotechnology, 2019, 17, 53.	9.1	15
121	Dark metabolism: a molecular insight into how the Antarctic seaâ€ice diatom <i>Fragilariopsis cylindrus</i> survives longâ€term darkness. New Phytologist, 2019, 223, 675-691.	7.3	40
122	Soil and Crop Management Practices to Minimize the Impact of Waterlogging on Crop Productivity. Frontiers in Plant Science, 2019, 10, 140.	3.6	120
123	Identification of QTL Related to ROS Formation under Hypoxia and Their Association with Waterlogging and Salt Tolerance in Barley. International Journal of Molecular Sciences, 2019, 20, 699.	4.1	42
124	Developing a high-throughput phenotyping method for oxidative stress tolerance in barley roots. Plant Methods, 2019, 15, 12.	4.3	16
125	Linking ploidy level with salinity tolerance: NADPH-dependent  ROS–Ca2+ hub' in the spotlight. Journal of Experimental Botany, 2019, 70, 1063-1067.	4.8	20
126	Friend or Foe? Chloride Patterning in Halophytes. Trends in Plant Science, 2019, 24, 142-151.	8.8	49

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127	Understanding physiological and morphological traits contributing to drought tolerance in barley. Journal of Agronomy and Crop Science, 2019, 205, 129-140.	3.5	34
128	Liver proteome response of pre-harvest Atlantic salmon following exposure to elevated temperature. BMC Genomics, 2018, 19, 133.	2.8	43
129	Fractionation of Dissolved Organic Matter on Coupled Reversed-Phase Monolithic Columns and Characterisation Using Reversed-Phase Liquid Chromatography-High Resolution Mass Spectrometry. Chromatographia, 2018, 81, 203-213.	1.3	11
130	Chronic β ₂ â€adrenoceptor agonist treatment alters muscle proteome and functional adaptations induced by high intensity training in young men. Journal of Physiology, 2018, 596, 231-252.	2.9	41
131	A multiple near isogenic line (multi-NIL) RNA-seq approach to identify candidate genes underpinning QTL. Theoretical and Applied Genetics, 2018, 131, 613-624.	3.6	30
132	Transcriptome and proteome profiling reveals stress-induced expression signatures of imiquimod-treated Tasmanian devil facial tumor disease (DFTD) cells. Oncotarget, 2018, 9, 15895-15914.	1.8	13
133	Understanding the Molecular Basis of Salt Sequestration in Epidermal Bladder Cells of Chenopodium quinoa. Current Biology, 2018, 28, 3075-3085.e7.	3.9	98
134	Proteomic analysis of Lactobacillus casei GCRL163 cell-free extracts reveals a SecB homolog and other biomarkers of prolonged heat stress. PLoS ONE, 2018, 13, e0206317.	2.5	15
135	Effects of exogenously-applied L-ascorbic acid on root expansive growth and viability of the border-like cells. Plant Signaling and Behavior, 2018, 13, e1514895.	2.4	5
136	Multi-lumen capillary based trypsin micro-reactor for the rapid digestion of proteins. Analyst, The, 2018, 143, 4944-4953.	3.5	11
137	Temporal changes in soil properties and physiological characteristics of Atriplex species and Medicago arborea grown in different soil types under saline irrigation. Plant and Soil, 2018, 432, 315-331.	3.7	4
138	Boron Alleviates Aluminum Toxicity by Promoting Root Alkalization in Transition Zone via Polar Auxin Transport. Plant Physiology, 2018, 177, 1254-1266.	4.8	65
139	Revealing mechanisms of salinity tissue tolerance in succulent halophytes: <scp>A</scp> case study for <scp><i>Carpobrotus rossi</i> Plant, Cell and Environment, 2018, 41, 2654-2667.</scp>	5.7	33
140	Towards complete identification of allergens in Jack Jumper (<i>Myrmecia pilosula</i>) ant venom and their clinical relevance: An immunoproteomic approach. Clinical and Experimental Allergy, 2018, 48, 1222-1234.	2.9	13
141	Heat shock proteins expressed in the marsupial Tasmanian devil are potential antigenic candidates in a vaccine against devil facial tumour disease. PLoS ONE, 2018, 13, e0196469.	2.5	6
142	An early ABA-induced stomatal closure, Na+ sequestration in leaf vein and K+ retention in mesophyll confer salt tissue tolerance in Cucurbita species. Journal of Experimental Botany, 2018, 69, 4945-4960.	4.8	77
143	Can highly saline irrigation water improve sodicity and alkalinity in sodic clayey subsoils?. Journal of Soils and Sediments, 2018, 18, 3290-3302.	3.0	7
144	It is not all about sodium: revealing tissue specificity and signalling roles of potassium in plant responses to salt stress. Plant and Soil, 2018, 431, 1-17.	3.7	245

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145	Piriformospora indica improves salinity stress tolerance in Zea mays L. plants by regulating Na+ and K+ loading in root and allocating K+ in shoot. Plant Growth Regulation, 2018, 86, 323-331.	3.4	71
146	Stomata in a saline world. Current Opinion in Plant Biology, 2018, 46, 87-95.	7.1	111
147	Calcium transport across plant membranes: mechanisms and functions. New Phytologist, 2018, 220, 49-69.	7.3	289
148	Reproductive Physiology of Halophytes: Current Standing. Frontiers in Plant Science, 2018, 9, 1954.	3.6	94
149	Agronomical, biochemical and histological response of resistant and susceptible wheat and barley under BYDV stress. Peerl, 2018, 6, e4833.	2.0	5
150	QTLs for stomatal and photosynthetic traits related to salinity tolerance in barley. BMC Genomics, 2017, 18, 9.	2.8	108
151	Triploid Atlantic salmon shows similar performance, fatty acid composition and proteome response to diploids during early freshwater rearing. Comparative Biochemistry and Physiology Part D: Genomics and Proteomics, 2017, 22, 67-77.	1.0	13
152	Chloroplast function and ion regulation in plants growing on saline soils: lessons from halophytes. Journal of Experimental Botany, 2017, 68, 3129-3143.	4.8	187
153	A high-quality genome assembly of quinoa provides insights into the molecular basis of salt bladder-based salinity tolerance and the exceptional nutritional value. Cell Research, 2017, 27, 1327-1340.	12.0	170
154	Signalling by potassium: another second messenger to add to the list?. Journal of Experimental Botany, 2017, 68, 4003-4007.	4.8	159
155	Genome-Wide Association Study Reveals a New QTL for Salinity Tolerance in Barley (Hordeum vulgare) Tj ETQq1	1 9.78431	.4 ggBT /Ove
156	Sequential protein extraction as an efficient method for improved proteome coverage in larvae of Atlantic salmon (Salmo salar). Proteomics, 2016, 16, 2043-2047.	2.2	4
157	Near-isogenic lines developed for a major QTL on chromosome arm 4HL conferring Fusarium crown rot resistance in barley. Euphytica, 2016, 209, 555-563.	1.2	21
158	Discovery of Biomarkers for Tasmanian Devil Cancer (DFTD) by Metabolic Profiling of Serum. Journal of Proteome Research, 2016, 15, 3827-3840.	3.7	13
159	Difference in root K ⁺ retention ability and reduced sensitivity of K ⁺ -permeable channels to reactive oxygen species confer differential salt tolerance in three <i>Brassica</i> species. Journal of Experimental Botany, 2016, 67, 4611-4625.	4.8	127
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