Richard Wilson

List of Publications by Year
in descending order

221 papers	10,168 citations	31976 53 h-index	46799 89 g-index
230 all docs	230 docs citations	230 times ranked	9417 citing authors

#	Article	IF	Citations
1	Learning from halophytes: physiological basis and strategies to improve abiotic stress tolerance in crops. Annals of Botany, 2013, 112, 1209-1221.	2.9	645
2	Mechanisms of Plant Responses and Adaptation to Soil Salinity. Innovation(China), 2020, 1, 100017.	9.1	387
3	Calcium transport across plant membranes: mechanisms and functions. New Phytologist, 2018, 220, 49-69.	7.3	289
4	Energy costs of salt tolerance in crop plants. New Phytologist, 2020, 225, 1072-1090.	7.3	284
5	Salt tolerance mechanisms in quinoa (Chenopodium quinoa Willd.). Environmental and Experimental Botany, 2013, 92, 43-54.	4.2	263
6	Salt bladders: do they matter?. Trends in Plant Science, 2014, 19, 687-691.	8.8	247
7	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
8	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
9	Salicylic acid in plant salinity stress signalling and tolerance. Plant Growth Regulation, 2015, 76, 25-40.	3.4	186
10	Oxidative stress protection and stomatal patterning as components of salinity tolerance mechanism in quinoa (Chenopodium quinoa). Physiologia Plantarum, 2012, 146, 26-38.	5.2	181
11	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
12	The translocation, folding, assembly and redox-dependent degradation of secretory and membrane proteins in semi-permeabilized mammalian cells. Biochemical Journal, 1995, 307, 679-687.	3.7	166
13	Signalling by potassium: another second messenger to add to the list?. Journal of Experimental Botany, 2017, 68, 4003-4007.	4.8	159
14	Doing â€˜business as usualâ€™ 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
15	Protein disulfide Isomerase Acts as a Molecular Chaperone during the Assembly of Procollagen. Journal of Biological Chemistry, 1998, 273, 9637-9643.	3.4	129
16	Non-stomatal limitation of photosynthesis by soil salinity. Critical Reviews in Environmental Science and Technology, 2021, 51, 791-825.	12.8	129
17	Cell surface and intracellular auxin signalling for H+ fluxes in root growth. Nature, 2021, 599, 273-277.	27.8	128
18	Amino acids regulate salinity-induced potassium efflux in barley root epidermis. Planta, 2007, 225, 753-761.	3.2	127

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19	Targeted Induction of Endoplasmic Reticulum Stress Induces Cartilage Pathology. PLoS Genetics, 2009, 5, e1000691.	3.5	127
20	Difference in root K⁺retention ability and reduced sensitivity of K⁺-permeable channels to reactive oxygen species confer differential salt tolerance in threeBrassicaspecies. Journal of Experimental Botany, 2016, 67, 4611-4625.	4.8	127
21	Soil and Crop Management Practices to Minimize the Impact of Waterlogging on Crop Productivity. Frontiers in Plant Science, 2019, 10, 140.	3.6	120
22	Ability of leaf mesophyll to retain potassium correlates with salinity tolerance in wheat and barley. Physiologia Plantarum, 2013, 149, 515-527.	5.2	113
23	Stomata in a saline world. Current Opinion in Plant Biology, 2018, 46, 87-95.	7.1	111
24	QTLs for stomatal and photosynthetic traits related to salinity tolerance in barley. BMC Genomics, 2017, 18, 9.	2.8	108
25	Understanding the Molecular Basis of Salt Sequestration in Epidermal Bladder Cells of Chenopodium quinoa. Current Biology, 2018, 28, 3075-3085.e7.	3.9	98
26	Metabolomics Data Normalization with EigenMS. PLoS ONE, 2014, 9, e116221.	2.5	96
27	Naxloci affect SOS1-like Na⁺/H⁺exchanger expression and activity in wheat. Journal of Experimental Botany, 2016, 67, 835-844.	4.8	95
28	Reproductive Physiology of Halophytes: Current Standing. Frontiers in Plant Science, 2018, 9, 1954.	3.6	94
29	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
30	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
31	Reducing Cadmium Accumulation in Plants: Structureâ€“Function Relations and Tissue-Specific Operation of Transporters in the Spotlight. Plants, 2020, 9, 223.	3.5	88
32	Extensive Gene Acquisition in the Extremely Psychrophilic Bacterial Species Psychroflexus torquis and the Link to Sea-Ice Ecosystem Specialism. Genome Biology and Evolution, 2014, 6, 133-148.	2.5	87
33	Linking salinity stress tolerance with tissue-specific Na+ sequestration in wheat roots. Frontiers in Plant Science, 2015, 6, 71.	3.6	86
34	The energy cost of the tonoplast futile sodium leak. New Phytologist, 2020, 225, 1105-1110.	7.3	86
35	Differentially expressed proteins in gill and skin mucus of Atlantic salmon (Salmo salar) affected by amoebic gill disease. Fish and Shellfish Immunology, 2014, 40, 69-77.	3.6	85
36	Genome-Wide Association Study Reveals a New QTL for Salinity Tolerance in Barley (Hordeum vulgare) Tj ETQq0 0 0 rgBT /Overlock 10 T	3.6	83

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37	Root vacuolar Na⁺ sequestration but not exclusion from uptake correlates with barley salt tolerance. Plant Journal, 2019, 100, 55-67.	5.7	80
38	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
39	Crop Halophytism: An Environmentally Sustainable Solution for Global Food Security. Trends in Plant Science, 2020, 25, 630-634.	8.8	77
40	Rutin, a flavonoid with antioxidant activity, improves plant salinity tolerance by regulating K+ retention and Na+ exclusion from leaf mesophyll in quinoa and broad beans. Functional Plant Biology, 2016, 43, 75.	2.1	76
41	Comprehensive Profiling of Cartilage Extracellular Matrix Formation and Maturation Using Sequential Extraction and Label-free Quantitative Proteomics. Molecular and Cellular Proteomics, 2010, 9, 1296-1313.	3.8	73
42	Changes in the Chondrocyte and Extracellular Matrix Proteome during Post-natal Mouse Cartilage Development. Molecular and Cellular Proteomics, 2012, 11, M111.014159.	3.8	73
43	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
44	GORK Channel: A Master Switch of Plant Metabolism?. Trends in Plant Science, 2020, 25, 434-445.	8.8	73
45	S100A8 and S100A9 in experimental osteoarthritis. Arthritis Research and Therapy, 2010, 12, R16.	3.5	72
46	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
47	Type-III procollagen assembly in semi-intact cells: chain association, nucleation and triple-helix folding do not require formation of inter-chain disulphide bonds but triple-helix nucleation does require hydroxylation. Biochemical Journal, 1996, 317, 195-202.	3.7	70
48	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
49	Boron Alleviates Aluminum Toxicity by Promoting Root Alkalization in Transition Zone via Polar Auxin Transport. Plant Physiology, 2018, 177, 1254-1266.	4.8	65
50	Mutations ofCOL10A1 in Schmid metaphyseal chondrodysplasia. Human Mutation, 2005, 25, 525-534.	2.5	64
51	AFB1 controls rapid auxin signalling through membrane depolarization in Arabidopsis thaliana root. Nature Plants, 2021, 7, 1229-1238.	9.3	59
52	Misfolding of Collagen X Chains Harboring Schmid Metaphyseal Chondrodysplasia Mutations Results in Aberrant Disulfide Bond Formation, Intracellular Retention, and Activation of the Unfolded Protein Response. Journal of Biological Chemistry, 2005, 280, 15544-15552.	3.4	58
53	Proteomic characterization of mouse cartilage degradation in vitro. Arthritis and Rheumatism, 2008, 58, 3120-3131.	6.7	58
54	Evaluating relative contribution of osmotolerance and tissue tolerance mechanisms toward salinity stress tolerance in three Brassica species. Physiologia Plantarum, 2016, 158, 135-151.	5.2	58

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55	Back to the Wild: On a Quest for Donors Toward Salinity Tolerant Rice. Frontiers in Plant Science, 2020, 11, 323.	3.6	54
56	Arsenic transport and interaction with plant metabolism: Clues for improving agricultural productivity and food safety. Environmental Pollution, 2021, 290, 117987.	7.5	54
57	Chromatographic methods for the isolation, separation and characterisation of dissolved organic matter. Environmental Sciences: Processes and Impacts, 2015, 17, 1531-1567.	3.5	52
58	Rewilding crops for climate resilience: economic analysis and de novo domestication strategies. Journal of Experimental Botany, 2021, 72, 6123-6139.	4.8	52
59	Light-stimulated growth of proteorhodopsin-bearing sea-ice psychrophile Psychroflexus torquis is salinity dependent. ISME Journal, 2013, 7, 2206-2213.	9.8	51
60	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
61	Friend or Foe? Chloride Patterning in Halophytes. Trends in Plant Science, 2019, 24, 142-151.	8.8	49
62	NADPH oxidases and the evolution of plant salinity tolerance. Plant, Cell and Environment, 2020, 43, 2957-2968.	5.7	49
63	Prospects for the accelerated improvement of the resilient crop quinoa. Journal of Experimental Botany, 2020, 71, 5333-5347.	4.8	49
64	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
65	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
66	Comparative Proteomic Analysis of Normal and Collagen IX Null Mouse Cartilage Reveals Altered Extracellular Matrix Composition and Novel Components of the Collagen IX Interactome. Journal of Biological Chemistry, 2013, 288, 13481-13492.	3.4	46
67	Improving Performance of Salt-Grown Crops by Exogenous Application of Plant Growth Regulators. Biomolecules, 2021, 11, 788.	4.0	46
68	Phylogenetic Diversity and Physiological Roles of Plant Monovalent Cation/H+ Antiporters. Frontiers in Plant Science, 2020, 11, 573564.	3.6	45
69	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
70	Fibrillin assembly: dimer formation mediated by amino-terminal sequences. Journal of Cell Science, 1999, 112, 3549-3558.	2.0	45
71	Liver proteome response of pre-harvest Atlantic salmon following exposure to elevated temperature. BMC Genomics, 2018, 19, 133.	2.8	43
72	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

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73	WARP Is a Novel Multimeric Component of the Chondrocyte Pericellular Matrix That Interacts with Perlecan. Journal of Biological Chemistry, 2006, 281, 7341-7349.	3.4	41
74	Chronic Î²₂â€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
75	Stomatal traits as a determinant of superior salinity tolerance in wild barley. Journal of Plant Physiology, 2020, 245, 153108.	3.5	41
76	Dark metabolism: a molecular insight into how the Antarctic seaâ€ice diatom Fragilariopsis cylindrus survives longâ€term darkness. New Phytologist, 2019, 223, 675-691.	7.3	40
77	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
78	Cartilage-specific ablation of XBP1 signaling in mouse results in a chondrodysplasia characterized by reduced chondrocyte proliferation and delayed cartilage maturation and mineralization. Osteoarthritis and Cartilage, 2015, 23, 661-670.	1.3	38
79	Impact of Lactose Starvation on the Physiology of Lactobacillus casei GCRL163 in the Presence or Absence of Tween 80. Journal of Proteome Research, 2013, 12, 5313-5322.	3.7	37
80	Identification of vitamin D sensitive pathways during lung development. Respiratory Research, 2016, 17, 47.	3.6	37
81	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
82	Metabolomics reveals increased isoleukotoxin diol (12,13-DHOME) in human plasma after acute Intralipid infusion. Journal of Lipid Research, 2012, 53, 1979-1986.	4.2	35
83	Understanding physiological and morphological traits contributing to drought tolerance in barley. Journal of Agronomy and Crop Science, 2019, 205, 129-140.	3.5	34
84	Calcium-Dependent Hydrogen Peroxide Mediates Hydrogen-Rich Water-Reduced Cadmium Uptake in Plant Roots. Plant Physiology, 2020, 183, 1331-1344.	4.8	34
85	A new allele for aluminium tolerance gene in barley (Hordeum vulgare L.). BMC Genomics, 2016, 17, 186.	2.8	33
86	Revealing mechanisms of salinity tissue tolerance in succulent halophytes: <scp>A</scp> case study for <scp>Carpobrotus rossi</scp>. Plant, Cell and Environment, 2018, 41, 2654-2667.	5.7	33
87	Biochemical pH clamp: the forgotten resource in membrane bioenergetics. New Phytologist, 2020, 225, 37-47.	7.3	33
88	Evolution of rapid blueâ€light response linked to explosive diversification of ferns in angiosperm forests. New Phytologist, 2021, 230, 1201-1213.	7.3	33
89	Proteomic analysis of mouse growth plate cartilage. Proteomics, 2006, 6, 6549-6553.	2.2	32
90	Proteomic analysis of cartilage proteins. Methods, 2008, 45, 22-31.	3.8	32

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91	Identification of new QTL for salt tolerance from rice variety Pokkali. Journal of Agronomy and Crop Science, 2020, 206, 202-213.	3.5	31
92	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
93	Collagen X Chains Harboring Schmid Metaphyseal Chondrodysplasia NC1 Domain Mutations Are Selectively Retained and Degraded in Stably Transfected Cells. Journal of Biological Chemistry, 2002, 277, 12516-12524.	3.4	29
94	Two of a kind: transmissible Schwann cell cancers in the endangered Tasmanian devil (Sarcophilus) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50	5.4	28
95	Lipid kinases PIP5K7 and PIP5K9 are required for polyamineâ€triggered K⁺ efflux in Arabidopsis roots. Plant Journal, 2020, 104, 416-432.	5.7	28
96	Novel Elements of the Chondrocyte Stress Response Identified Using an in Vitro Model of Mouse Cartilage Degradation. Journal of Proteome Research, 2016, 15, 1033-1050.	3.7	27
97	Melatonin as a regulator of plant ionic homeostasis: implications for abiotic stress tolerance. Journal of Experimental Botany, 2022, 73, 5886-5902.	4.8	26
98	Antioxidant Enzymatic Activity and Osmotic Adjustment as Components of the Drought Tolerance Mechanism in Carex duriuscula. Plants, 2021, 10, 436.	3.5	25
99	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
100	Quantitative Trait Loci for Salinity Tolerance Identified under Drained and Waterlogged Conditions and Their Association with Flowering Time in Barley (Hordeum vulgare. L). PLoS ONE, 2015, 10, e0134822.	2.5	25
101	Proteomics makes progress in cartilage and arthritis research. Matrix Biology, 2009, 28, 121-128.	3.6	24
102	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
103	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
104	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
105	Function of NHX-type transporters in improving rice tolerance to aluminum stress and soil acidity. Planta, 2020, 251, 71.	3.2	23
106	Rewilding staple crops for the lost halophytism: Toward sustainability and profitability of agricultural production systems. Molecular Plant, 2022, 15, 45-64.	8.3	23
107	Comparing Kinetics of Xylem Ion Loading and Its Regulation in Halophytes and Glycophytes. Plant and Cell Physiology, 2020, 61, 403-415.	3.1	22
108	Sodium sequestration confers salinity tolerance in an ancestral wild rice. Physiologia Plantarum, 2021, 172, 1594-1608.	5.2	22

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109	Tissue tolerance mechanisms conferring salinity tolerance in a halophytic perennial species Nitraria sibirica Pall.. Tree Physiology, 2021, 41, 1264-1277.	3.1	22
110	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
111	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
112	A robust method for proteomic characterization of mouse cartilage using solubility-based sequential fractionation and two-dimensional gel electrophoresis. Matrix Biology, 2008, 27, 709-712.	3.6	20
113	Linking ploidy level with salinity tolerance: NADPH-dependent â€˜ROSâ€“Ca2+ hubâ€™ in the spotlight. Journal of Experimental Botany, 2019, 70, 1063-1067.	4.8	20
114	Salinity Effects on Guard Cell Proteome in Chenopodium quinoa. International Journal of Molecular Sciences, 2021, 22, 428.	4.1	20
115	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
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	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
118	A hemeâ€bbinding protein produced by Haemophilus haemolyticus inhibits nonâ€typeable Haemophilus influenzae. Molecular Microbiology, 2020, 113, 381-398.	2.5	18
119	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
120	Understanding the mechanistic basis of adaptation of perennial Sarcocornia quinqueflora species to soil salinity. Physiologia Plantarum, 2021, 172, 1997-2010.	5.2	18
121	Shotgun Proteomics as a Powerful Tool for the Study of the Proteomes of Plants, Their Pathogens, and Plantâ€“Pathogen Interactions. Proteomes, 2022, 10, 5.	3.5	18
122	The extracellular matrix: an underexplored but important proteome. Expert Review of Proteomics, 2010, 7, 803-806.	3.0	17
123	Enhancing Fusarium crown rot resistance by pyramiding large-effect QTL in barley. Molecular Breeding, 2015, 35, 1.	2.1	16
124	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
125	Developing a high-throughput phenotyping method for oxidative stress tolerance in barley roots. Plant Methods, 2019, 15, 12.	4.3	16
126	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

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127	Sugar Beet (Beta vulgaris) Guard Cells Responses to Salinity Stress: A Proteomic Analysis. International Journal of Molecular Sciences, 2020, 21, 2331.	4.1	16
128	Understanding a Mechanistic Basis of ABA Involvement in Plant Adaptation to Soil Flooding: The Current Standing. Plants, 2021, 10, 1982.	3.5	16
129	Mitochondrial respiratory chain function promotes extracellular matrix integrity in cartilage. Journal of Biological Chemistry, 2021, 297, 101224.	3.4	16
130	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
131	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
132	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
133	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
134	Evaluation of salt tolerance of oat cultivars and the mechanism of adaptation to salinity. Journal of Plant Physiology, 2022, 273, 153708.	3.5	15
135	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
136	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
137	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
138	Attached and Planktonic Listeria monocytogenes Global Proteomic Responses and Associated Influence of Strain Genetics and Temperature. Journal of Proteome Research, 2015, 14, 1161-1173.	3.7	13
139	Discovery of Biomarkers for Tasmanian Devil Cancer (DFTD) by Metabolic Profiling of Serum. Journal of Proteome Research, 2016, 15, 3827-3840.	3.7	13
140	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
141	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
142	Towards complete identification of allergens in Jack Jumper (Myrmecia pilosula) ant venom and their clinical relevance: An immunoproteomic approach. Clinical and Experimental Allergy, 2018, 48, 1222-1234.	2.9	13
143	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
144	Plant responses to heterogeneous salinity: agronomic relevance and research priorities. Annals of Botany, 2022, 129, 499-518.	2.9	13

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145	Proteomic Insight into Functional Changes of Proteorhodopsin-Containing Bacterial Species Psychroflexus torquis under Different Illumination and Salinity Levels. Journal of Proteome Research, 2015, 14, 3848-3858.	3.7	12
146	A Disintegrin and Metalloproteinase with Thrombospondin Motifs-5 (ADAMTS-5) Forms Catalytically Active Oligomers. Journal of Biological Chemistry, 2016, 291, 3197-3208.	3.4	12
147	The metabolomics of alpha-synuclein (SNCA) gene deletion and mutation in mouse brain. Metabolomics, 2014, 10, 114-122.	3.0	11
148	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
149	Multi-lumen capillary based trypsin micro-reactor for the rapid digestion of proteins. Analyst, The, 2018, 143, 4944-4953.	3.5	11
150	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
151	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
152	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
153	Early responses to salt stress in quinoa genotypes with opposite behavior. Physiologia Plantarum, 2021, 173, 1392-1420.	5.2	10
154	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
155	An RNA-binding protein MUG13.4 interacts with AtAGO2 to modulate salinity tolerance in Arabidopsis. Plant Science, 2019, 288, 110218.	3.6	9
156	Optimisation of Sporosori Purification and Protein Extraction Techniques for the Biotrophic Protozoan Plant Pathogen Spongospora subterranea. Molecules, 2020, 25, 3109.	3.8	9
157	Quantitative proteomics provides an insight into germinationâ€related proteins in the obligate biotrophic plant pathogen <scp>Spongospora subterranea</scp>. Environmental Microbiology Reports, 2021, 13, 521-532.	2.4	9
158	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
159	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
160	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
161	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
162	Local and systemic responses conferring acclimation of Brassica napus roots to low phosphorus conditions. Journal of Experimental Botany, 2022, 73, 4753-4777.	4.8	9

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163	Cartilage proteomics: Challenges, solutions and recent advances. Proteomics - Clinical Applications, 2008, 2, 251-263.	1.6	8
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