

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Comparative transcriptomic analysis uncovers conserved pathways involved in adventitious root formation in poplar. Physiology and Molecular Biology of Plants, 2021, 27, 1903-1918.	3.1	10
2	Pretreating poplar cuttings with low nitrogen ameliorates salt stress responses by increasing stored carbohydrates and priming stress signaling pathways. Ecotoxicology and Environmental Safety, 2021, 225, 112801.	6.0	8
3	Integrating multiple omics to identify common and specific molecular changes occurring in Arabidopsis under chronic nitrate and sulfate limitations. Journal of Experimental Botany, 2020, 71, 6471-6490.	4.8	18
4	Integrated Transcriptome Analysis Reveals Plant Hormones Jasmonic Acid and Salicylic Acid Coordinate Growth and Defense Responses upon Fungal Infection in Poplar. Biomolecules, 2019, 9, 12.	4.0	72
5	Evolutionary analyses of NIN-like proteins in plants and their roles in nitrate signaling. Cellular and Molecular Life Sciences, 2019, 76, 3753-3764.	5.4	67
6	Proteomic and lipidomic analyses of the Arabidopsis <i>atg5</i> autophagy mutant reveal major changes in endoplasmic reticulum and peroxisome metabolisms and in lipid composition. New Phytologist, 2019, 223, 1461-1477.	7.3	54
7	Global Transcriptomic Profile Analysis of Genes Involved in Lignin Biosynthesis and Accumulation Induced by Boron Deficiency in Poplar Roots. Biomolecules, 2019, 9, 156.	4.0	19
8	Growth performance, photosynthesis, and root characteristics are associated with nitrogen use efficiency in six poplar species. Environmental and Experimental Botany, 2019, 164, 40-51.	4.2	28
9	Morphological and physiological responses to contrasting nitrogen regimes in Populus cathayana is linked to resources allocation and carbon/nitrogen partition. Environmental and Experimental Botany, 2019, 162, 247-255.	4.2	45
10	Genome-wide identification of BOR genes in poplar and their roles in response to various environmental stimuli. Environmental and Experimental Botany, 2019, 164, 101-113.	4.2	16
11	Autophagy and Nutrients Management in Plants. Cells, 2019, 8, 1426.	4.1	50
12	The PIN-FORMED Auxin Efflux Carriers in Plants. International Journal of Molecular Sciences, 2018, 19, 2759.	4.1	113
13	Aux/IAA Gene Family in Plants: Molecular Structure, Regulation, and Function. International Journal of Molecular Sciences, 2018, 19, 259.	4.1	277
14	Comparative transcriptomic analysis reveals the roles of overlapping heat-/drought-responsive genes in poplars exposed to high temperature and drought. Scientific Reports, 2017, 7, 43215.	3.3	72
15	Uncovering the physiological mechanisms that allow nitrogen availability to affect drought acclimation in Catalpa bungei. Tree Physiology, 2017, 37, 1453-1456.	3.1	10
16	Identification of TIFY Family Genes and Analysis of Their Expression Profiles in Response to Phytohormone Treatments and Melampsora larici-populina Infection in Poplar. Frontiers in Plant Science, 2017, 8, 493.	3.6	33
17	The conserved salt-responsive genes in the roots of Populus×canescens and Arabidopsis thaliana. Environmental and Experimental Botany, 2016, 129, 48-56.	4.2	23
18	Global poplar root and leaf transcriptomes reveal links between growth and stress responses under nitrogen starvation and excess. Tree Physiology, 2015, 35, 1283-1302.	3.1	131

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19	Ectomycorrhizas with <i><scp>P</scp>axillus involutus</i> enhance cadmium uptake and tolerance in <i><scp>P</scp>opulus</i> × <i>canescens</i> . Plant, Cell and Environment, 2014, 37, 627-642.	5.7	118
20	Net fluxes of ammonium and nitrate in association with H+ fluxes in fine roots of Populus popularis. Planta, 2013, 237, 919-931.	3.2	112
21	Nitrogen metabolism of two contrasting poplar species during acclimation to limiting nitrogen availability. Journal of Experimental Botany, 2013, 64, 4207-4224.	4.8	180
22	N-fertilization has different effects on the growth, carbon and nitrogen physiology, and wood properties of slow- and fast-growing Populus species. Journal of Experimental Botany, 2012, 63, 6173-6185.	4.8	131