Karolina Chwialkowska

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/1462994/publications.pdf

Version: 2024-02-01

20 papers 822 citations

759055 12 h-index 19 g-index

20 all docs

20 docs citations

times ranked

20

1361 citing authors

#	Article	IF	CITATIONS
1	Molecular Signature of Subtypes of Non-Small-Cell Lung Cancer by Large-Scale Transcriptional Profiling: Identification of Key Modules and Genes by Weighted Gene Co-Expression Network Analysis (WGCNA). Cancers, 2020, 12, 37.	1.7	179
2	A multi-tissue full lifespan epigenetic clock for mice. Aging, 2018, 10, 2832-2854.	1.4	166
3	Ancient Evolutionary Trade-Offs between Yeast Ploidy States. PLoS Genetics, 2013, 9, e1003388.	1.5	85
4	Transcriptome analysis reveals the role of the root hairs as environmental sensors to maintain plant functions under water-deficiency conditions. Journal of Experimental Botany, 2016, 67, 1079-1094.	2.4	80
5	Water-deficiency conditions differently modulate the methylome of roots and leaves in barley (<i>Hordeum vulgare</i> L.). Journal of Experimental Botany, 2016, 67, 1109-1121.	2.4	72
6	Accumulation of peroxidase-related reactive oxygen species in trichoblasts correlates with root hair initiation in barley. Journal of Plant Physiology, 2013, 170, 185-195.	1.6	45
7	Methylation Sensitive Amplification Polymorphism Sequencing (MSAP-Seq)—A Method for High-Throughput Analysis of Differentially Methylated CCGG Sites in Plants with Large Genomes. Frontiers in Plant Science, 2017, 8, 2056.	1.7	32
8	DNA Demethylation in Response to Heat Stress in Arabidopsis thaliana. International Journal of Molecular Sciences, 2021, 22, 1555.	1.8	31
9	iRootHair: A Comprehensive Root Hair Genomics Database Â. Plant Physiology, 2012, 161, 28-35.	2.3	30
10	Molecular changes in vitellogenin gene of Spodoptera exigua after long-time exposure to cadmium – Toxic side effect or microevolution?. Ecotoxicology and Environmental Safety, 2018, 147, 461-470.	2.9	21
11	Aluminum or Low pH – Which Is the Bigger Enemy of Barley? Transcriptome Analysis of Barley Root Meristem Under Al and Low pH Stress. Frontiers in Genetics, 2021, 12, 675260.	1.1	21
12	A comparative analysis of proteins that accumulate during the initial stage of root hair development in barley root hair mutants and their parent varieties. Journal of Applied Genetics, 2012, 53, 363-376.	1.0	17
13	Insights into the Histone Acetylation-Mediated Regulation of the Transcription Factor Genes That Control the Embryogenic Transition in the Somatic Cells of Arabidopsis. Cells, 2022, 11, 863.	1.8	13
14	Response of rhizospheric and endophytic bacterial communities of white mustard (Sinapis alba) to bioaugmentation of soil with the Pseudomonas sp. H15 strain. Ecotoxicology and Environmental Safety, 2020, 194, 110434.	2.9	10
15	Isolation and Characterization of Simple Sequence Repeats (SSR) Markers from the Moss Genus Orthotrichum Using a Small Throughput Pyrosequencing Machine. International Journal of Molecular Sciences, 2012, 13, 7586-7593.	1.8	9
16	Isolation and characterization of microsatellite loci in Pulsatilla patens (L.) Mill. (Ranunculaceae) a rare and endangered plant species in Europe. Conservation Genetics Resources, 2013, 5, 421-423.	0.4	6
17	Development of Microsatellite Markers Using Pyrosequencing in Galium trifidum (Rubiaceae), a Rare Species in Central Europe. International Journal of Molecular Sciences, 2012, 13, 9893-9899.	1.8	3
18	DNA Methylation Analysis in Barley and Other Species with Large Genomes. Methods in Molecular Biology, 2019, 1900, 253-268.	0.4	1

#	Article	IF	CITATIONS
19	Do Not Miss the (Genetic) Diagnosis of Gaucher Syndrome: A Narrative Review on Diagnostic Clues and Management in Severe Prenatal and Perinatal-Lethal Sporadic Cases. Journal of Clinical Medicine, 2021, 10, 4890.	1.0	1
20	Automatic analysis of 2D polyacrylamide gels in the diagnosis of DNA polymorphisms. BioMedical Engineering OnLine, 2013, 12, 68.	1.3	0