Anastasiya Krinitsyna

List of Publications by Year
in descending order

23 papers	409 citations	840776 11 h-index	752698 20 g-index
23 all docs	23 docs citations	23 times ranked	456 citing authors

#	Article	IF	Citations
1	Phylogeny, biogeography and systematics of Dysphanieae (Amaranthaceae). Taxon, 2021, 70, 526-551.	0.7	6
2	Scorzonera sensu lato (Asteraceae, Cichorieae) â€“ taxonomic reassessment in the light of new molecular phylogenetic and carpological analyses. PhytoKeys, 2020, 137, 1-85.	1.0	18
3	Molecular phylogenetic data and seed coat anatomy resolve the generic position of some critical Chenopodioideae (Chenopodiaceae â€“ Amaranthaceae) with reduced perianth segments. PhytoKeys, 2018, 109, 103-128.	1.0	15
4	Seed characters in Molluginaceae (Caryophyllales): implications for taxonomy and evolution. Botanical Journal of the Linnean Society, 2018, 187, 167-208.	1.6	8
5	Diagnostics, taxonomy, nomenclature and distribution of perennial Sesuvium (Aizoaceae) in Africa. PhytoKeys, 2018, 92, 45-88.	1.0	10
6	Comparative analysis of inverted repeats of polypod fern (Polypodiales) plastomes reveals two hypervariable regions. BMC Plant Biology, 2017, 17, 255.	3.6	23
7	miR319, miR390, and miR393 Are Involved in Aluminum Response in Flax (Linum usitatissimum L.). BioMed Research International, 2017, 2017, 1-6.	1.9	26
8	The systematic position of Dryopteris blanfordii subsp. nigrosquamosa (Ching) Fraser-Jenkins within the genus Dryopteris Adans.. PhytoKeys, 2017, 90, 89-112.	1.0	1
9	Identification, Expression Analysis, and Target Prediction of Flax Genotroph MicroRNAs Under Normal and Nutrient Stress Conditions. Frontiers in Plant Science, 2016, 7, 399.	3.6	43
10	Gene expression profiling of flax (Linum usitatissimum L.) under edaphic stress. BMC Plant Biology, 2016, 16, 237.	3.6	68
11	One-Seeded Fruits in the Core Caryophyllales: Their Origin and Structural Diversity. PLoS ONE, 2015, 10, e0117974.	2.5	36
12	A rapid and cost-effective method for DNA extraction from archival herbarium specimens. Biochemistry (Moscow), 2015, 80, 1478-1484.	1.5	18
13	Excess fertilizer responsive miRNAs revealed in Linum usitatissimum L. Biochimie, 2015, 109, 36-41.	2.6	31
14	Retrotransposon-Based Molecular Markers for Analysis of Genetic Diversity within the GenusLinum. BioMed Research International, 2014, 2014, 1-14.	1.9	24
15	Polymorphism of the KPI-A gene sequence in the potato subgenera Potatoe (Sect. Petota, Esolonifera,) Tj ETQq1 1 0.784314 rgBT /Ov	1.3	0
16	New combinations in Asiatic Oxybasis (Amaranthaceae s.l.): evidence from morphological, carpological and molecular data. Phytotaxa, 2013, 144, 1.	0.3	9
17	Impact of recombination on polymorphism of genes encoding Kunitz-type protease inhibitors in the genus Solanum. Biochimie, 2012, 94, 1687-1696.	2.6	9
18	Polygalacturonase-inhibiting protein is a structural component of plant cell wall. Biochemistry (Moscow), 2008, 73, 1053-1062.	1.5	32

#	Article	IF	Citations
19	Heterologous expression, purification, and properties of a potato protein inhibitor of serine proteinases. Biochemistry (Moscow), 2006, 71, 1176-1182.	1.5	9
20	Cloning of polygalacturonase inhibitor protein genes from Solanum brevidens Fill.. Russian Journal of Genetics, 2006, 42, 376-384.	0.6	1
21	Activity of polygalacturonase-inhibiting protein from banana fruit tissues. Applied Biochemistry and Microbiology, 2005, 41, 251-253.	0.9	7
22	Molecular cloning of Kunitz-type proteinase inhibitor group B genes from potato. Biochemistry (Moscow), 2005, 70, 292-299.	1.5	7
23	Role of the Polygalacturonidase Inhibitor Protein in the Ripening of Apples and Their Resistance to Monilia fructigena, a Causative Agent of Fruit Rot. Applied Biochemistry and Microbiology, 2004, 40, 89-92.	0.9	8