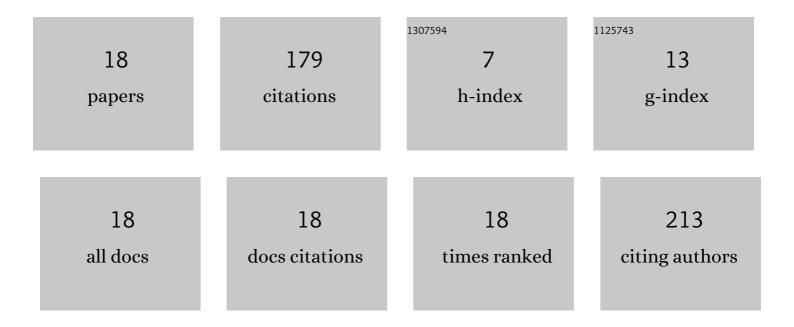
## Piotr Androsiuk

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3281937/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	RAPID ENVIRONMENTAL CHANGES IN THE WESTERN ANTARCTIC PENINSULA REGION DUE TO CLIMATE CHANGE AND HUMAN ACTIVITY. Applied Ecology and Environmental Research, 2017, 15, 525-539.	0.5	34
2	Genetic status of Norway spruce (Picea abies) breeding populations for northern Sweden. Silvae Genetica, 2013, 62, 127-136.	0.8	25
3	Genetic variability of Colobanthus quitensis from King George Island (Antarctica). Polish Polar Research, 2015, 36, 281-295.	0.9	20
4	Evolutionary dynamics of the chloroplast genome sequences of six Colobanthus species. Scientific Reports, 2020, 10, 11522.	3.3	19
5	Range-wide pattern of genetic variation in Colobanthus quitensis. Polar Biology, 2018, 41, 2467-2479.	1.2	16
6	How much of the invader's genetic variability can slip between our fingers? A case study of secondary dispersal of Poa annua on King George Island (Antarctica). Ecology and Evolution, 2018, 8, 592-600.	1.9	14
7	Retrotransposonâ€based genetic diversity of <i>Deschampsia antarctica</i> Desv. from King George Island (Maritime Antarctic). Ecology and Evolution, 2021, 11, 648-663.	1.9	9
8	The complete chloroplast genome of <i>Colobanthus apetalus</i> (Labill.) Druce: genome organization and comparison with related species. PeerJ, 2018, 6, e4723.	2.0	9
9	Triticale Green Plant Regeneration Is Due to DNA Methylation and Sequence Changes Affecting Distinct Sequence Contexts in the Presence of Copper Ions in Induction Medium. Cells, 2022, 11, 84.	4.1	7
10	Genetic variability of Pinus sylvestris populations from IUFRO 1982 provenance trial. Dendrobiology, 0, 71, 23-33.	0.6	6
11	Retrotransposon-based genetic variation of <i>Poa annua</i> populations from contrasting climate conditions. PeerJ, 2019, 7, e6888.	2.0	5
12	rps3 as a Candidate Mitochondrial Gene for the Molecular Identification of Species from the Colletotrichum acutatum Species Complex. Genes, 2020, 11, 552.	2.4	4
13	Characterization and phylogenetic analysis of the complete mitochondrial genome of the pathogenic fungus llyonectria destructans. Scientific Reports, 2022, 12, 2359.	3.3	4
14	Genetic diversity and differentiation of Pinus sylvestris L. from the IUFRO 1982 provenance trial revealed by AFLP analysis. Archives of Biological Sciences, 2015, 67, 1237-1249.	0.5	3
15	B-SAP markers derived from the bacterial KatG gene differentiate populations of Pinus sylvestris and provide new insights into their postglacial history. Silva Fennica, 2011, 45, .	1.3	2
16	Molecular Diversity and Phylogeny Reconstruction of Genus Colobanthus (Caryophyllaceae) Based on Mitochondrial Gene Sequences. Genes, 2022, 13, 1060.	2.4	2
17	Genetic and phenotypic relationships among Pinus sylvestris populations in the Pieniny National Park. Archives of Biological Sciences, 2018, 70, 289-297.	0.5	0
18	The effects of methanesulfonic acid on seed germination and morphophysiological changes in the seedlings of two Colobanthus species. Acta Societatis Botanicorum Poloniae, 2018, 87, .	0.8	0