Anastasiia K Kimeklis

List of Publications by Year
in descending order

19	206	1163117 8	1058476 14
papers	citations	h-index	g-index

25	25	25	257
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Bosea vaviloviae sp. nov., a new species of slow-growing rhizobia isolated from nodules of the relict species Vavilovia formosa (Stev.) Fed.. Antonie Van Leeuwenhoek, 2015, 107, 911-920.	1.7	51
2	Relationships between pasture legumes, rhizobacteria and nodule bacteria in heavy metal polluted mine waste of SW Sardinia. Symbiosis, 2012, 58, 149-159.	2.3	30
3	Genetic diversity of rhizobia isolated from nodules of the relic species Vavilovia formosa (Stev.) Fed.. Antonie Van Leeuwenhoek, 2014, 105, 389-399.	1.7	28
4	Extra-slow-growing Tardiphaga strains isolated from nodules of Vavilovia formosa (Stev.) Fed.. Archives of Microbiology, 2015, 197, 889-898.	2.2	15
5	Soil microbiome of the postmining areas in polar ecosystems in surroundings of Nadym, Western Siberia, Russia. Open Agriculture, 2019, 4, 684-696.	1.7	13
6	Characteristics of natural selection in populations of nodule bacteria (Rhizobium leguminosarum) interacting with different host plants. Russian Journal of Genetics, 2015, 51, 949-956.	0.6	11
7	Evolution of fixNOQP genes encoding cytochrome oxidase with high affinity to oxygen in rhizobia and related bacteria. Russian Journal of Genetics, 2017, 53, 766-774.	0.6	10
8	Rhizobia Isolated from the Relict Legume Vavilovia formosa Represent a Genetically Specific Group within Rhizobium leguminosarum biovar viciae. Genes, 2019, 10, 991.	2.4	10
9	Search for Ancestral Features in Genomes of Rhizobium leguminosarum bv. viciae Strains Isolated from the Relict Legume Vavilovia formosa. Genes, 2019, 10, 990.	2.4	8
10	The difference between cellulolytic â€˜culturomesâ€™ and microbiomes inhabiting two contrasting soil types. PLoS ONE, 2020, 15, e0242060.	2.5	6
11	Divergent Evolution of Symbiotic Bacteria: Rhizobia of the Relic Legume Vavilovia formosa Form an Isolated Group within Rhizobium leguminosarum bv. viciae. Russian Journal of Genetics, 2018, 54, 866-870.	0.6	5
12	Restoration of soil microbiome in various soil horizons after crown and surface wildfires. Ecological Genetics, 2020, 18, 343-356.	0.5	5
13	ANALYSIS OF MICROBIOME OF RECULTIVATED SOILS OF THE KINGISEPP AREA OF PHOSPHORITE MINING. Sel'skokhozyaistvennaya Biologiya, 2020, 55, 137-152.	0.3	4
14	The plastid and mitochondrial genomes of <i>Vavilovia Formosa</i> (Stev.) Fed. and the phylogeny of related legume genera. Vavilovskii Zhurnal Genetiki I Selektsii, 2020, 23, 972-980.	1.1	4
15	Structural and functional organization of the plasmid regulons of Rhizobium leguminosarum symbiotic genes. Microbiology, 2016, 85, 708-716.	1.2	2
16	Microbiomes of different ages in Rendzic Leptosols in the Crimean Peninsula. PeerJ, 2021, 9, e10871.	2.0	2
17	PHYLOGENETIC ANALYSIS OF Rhizobium STRAINS, ISOLATED FROM NODULES OF Vavilovia formosa (Stev.) Fed.. Sel'skokhozyaistvennaya Biologiya, 2015, 50, 655-664.	0.3	2
18	Evolutionary Geography of Root Nodule Bacteria: Speciation Directed by the Host Plants. Microbiology, 2020, 89, 1-12.	1.2	0

#	Article	IF	Citations
19	Microbiome composition of disturbed soils from sandy-gravel mining complexes with different reclamation approaches. One Ecosystem, 0, 7, .	0.0	0