## PaweÅ, Cyplik

List of Publications by Year in descending order

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430442 454577 34 945 18 30 citations h-index g-index papers 34 34 34 1182 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	The Influence of Bacteria Causing Subclinical Mastitis on the Structure of the Cow's Milk Microbiome. Molecules, 2022, 27, 1829.	1.7	18
2	Effect of Processing Treatment and Modified Atmosphere Packing on Carrot's Microbial Community Structure by Illumina MiSeq Sequencing. Molecules, 2022, 27, 2830.	1.7	1
3	Dose–Response Effect of Nitrogen on Microbial Community during Hydrocarbon Biodegradation in Simplified Model System. Applied Sciences (Switzerland), 2022, 12, 6012.	1.3	5
4	The Raw Milk Microbiota from Semi-Subsistence Farms Characteristics by NGS Analysis Method. Molecules, 2021, 26, 5029.	1.7	16
5	Heavy Metals as a Factor Increasing the Functional Genetic Potential of Bacterial Community for Polycyclic Aromatic Hydrocarbon Biodegradation. Molecules, 2020, 25, 319.	1.7	17
6	Assessment of soil potential to natural attenuation and autochthonous bioaugmentation using microarray and functional predictions from metagenome profiling. Annals of Microbiology, 2019, 69, 945-955.	1.1	4
7	The impact of natural and synthetic surfactants on bacterial community during hydrocarbon biodegradation. International Biodeterioration and Biodegradation, 2019, 142, 191-199.	1.9	14
8	The Toxic Effect of Herbicidal Ionic Liquids on Biogas-Producing Microbial Community. International Journal of Environmental Research and Public Health, 2019, 16, 916.	1.2	6
9	Multidimensional Toxicity of Rhamnolipid Extracts Obtained From Creosote-Contaminated Soil. Clean - Soil, Air, Water, 2018, 46, 1800053.	0.7	6
10	Characterization of specific spoilage organisms (SSOs) in vacuum-packed ham by culture-plating techniques and MiSeq next-generation sequencing technologies. Journal of the Science of Food and Agriculture, 2017, 97, 659-668.	1.7	13
11	Influence of soil contamination with PAH on microbial community dynamics and expression level of genes responsible for biodegradation of PAH and production of rhamnolipids. Environmental Science and Pollution Research, 2016, 23, 23043-23056.	2.7	35
12	Evaluating robustness of a diesel-degrading bacterial consortium isolated from contaminated soil. New Biotechnology, 2016, 33, 852-859.	2.4	30
13	Methane fermentation of the maize straw silage under meso- and thermophilic conditions. Energy, 2016, 115, 1495-1502.	4.5	66
14	Influence of oligomeric herbicidal ionic liquids with MCPA and Dicamba anions on the community structure of autochthonic bacteria present in agricultural soil. Science of the Total Environment, 2016, 563-564, 247-255.	3.9	49
15	Antibacterial effect of the Trichoderma viride fungi on soil microbiome during PAH's biodegradation. International Biodeterioration and Biodegradation, 2015, 104, 170-177.	1.9	27
16	High Voltage Electrochemiluminescence (ECL) as a New Method for Detection of PAH During Screening for PAH-Degrading Microbial Consortia. Water, Air, and Soil Pollution, 2015, 226, 270.	1.1	2
17	Removal of nitrates from processing wastewater by cryoconcentration combined with biological denitrification. Desalination and Water Treatment, 2015, 54, 1903-1911.	1.0	2
18	Biodegradation of diesel/biodiesel blends in saturated sand microcosms. Fuel, 2014, 116, 321-327.	3.4	58

#	Article	IF	CITATIONS
19	Biodegradation of Triton X-100 and its primary metabolites by a bacterial community isolated from activated sludge. Journal of Environmental Management, 2013, 128, 292-299.	3.8	24
20	Composting of oiled bleaching earth: Fatty acids degradation, phytotoxicity and mutagenicity changes. International Biodeterioration and Biodegradation, 2013, 78, 49-57.	1.9	43
21	Denitrification of industrial wastewater: Influence of glycerol addition on metabolic activity and community shifts in a microbial consortium. Chemosphere, 2013, 93, 2823-2831.	4.2	25
22	Biological denitrification of brine: the effect of compatible solutes on enzyme activities and fatty acid degradation. Biodegradation, 2012, 23, 663-672.	1.5	14
23	Rhamnolipids Increase the Phytotoxicity of Diesel Oil Towards Four Common Plant Species in a Terrestrial Environment. Water, Air, and Soil Pollution, 2012, 223, 4275-4282.	1.1	32
24	Biological Denitrification of High Nitrate Processing Wastewaters from Explosives Production Plant. Water, Air, and Soil Pollution, 2012, 223, 1791-1800.	1.1	38
25	Biodegradation of rhamnolipids in liquid cultures: Effect of biosurfactant dissipation on diesel fuel/B20 blend biodegradation efficiency and bacterial community composition. Bioresource Technology, 2012, 111, 328-335.	4.8	73
26	Genetic and chemical analyzes of transformations in compost compounds during biodegradation of oiled bleaching earth with waste sludge. Bioresource Technology, 2012, 114, 75-83.	4.8	5
27	Utilization of Triton X-100 and polyethylene glycols during surfactant-mediated biodegradation of diesel fuel. Journal of Hazardous Materials, 2011, 197, 97-103.	6.5	32
28	Relative quantitative PCR to assess bacterial community dynamics during biodegradation of diesel and biodiesel fuels under various aeration conditions. Bioresource Technology, 2011, 102, 4347-4352.	4.8	54
29	In Vitro Studies on Atrazine Effects on Human Intestinal Cells. Water, Air, and Soil Pollution, 2010, 213, 401-411.	1.1	15
30	Biodegradation and surfactant-mediated biodegradation of diesel fuel by 218 microbial consortia are not correlated to cell surface hydrophobicity. Applied Microbiology and Biotechnology, 2009, 84, 545-553.	1.7	79
31	The kinetics of nicotine degradation, enzyme activities and genotoxic potential in the characterization of tobacco waste composting. Bioresource Technology, 2009, 100, 5037-5044.	4.8	60
32	Atrazine degradation by aerobic microorganisms isolated from the rhizosphere of sweet flag (Acorus) Tj ETQq0	0 0 fgBT /e	Overlock 10 Tf
33	Application of a membrane bioreactor to denitrification of brine. Desalination, 2007, 207, 134-143.	4.0	28
34	Effect of macro/micro nutrients and carbon source over the denitrification rate of Haloferax denitrificans archaeon. Enzyme and Microbial Technology, 2007, 40, 212-220.	1.6	13