### **Decomposition and Carbon Cycling**

- 90. Tobias Arnstadt Log decay of *Fagus sylvatica* in temperate forests and the significance of lignin modifying enzymes for the degradation process
- 91. Chris Bamminger Divergent effects of pyrochar and hydrochar on greenhouse gas emissions and microbial abundances in an arable soil
- 92. Andrea Burešová Composition and activity of microbial community during decomposition of plant litter on two contrasting localities
- 93. Juliana Conceição The management system can influence the physiological function and social interaction of phosphate solubilizing bacteria isolated rhizosphere of *Carica papaya* L.
- 94. Ivana Eichlerová Decomposition traits and enzyme production of saprotrophic fungi are shaped by the combination of their ecophysiology and taxonomy
- 95. Lia del Pilar Fernández Soil bacterial diversity from different animal settlements in maritime Antarctica
- 96. Damien Finn Carbon and nitrogen co-metabolism and microbial nitrogen-mining both determine the extent of plant material decomposition in four Australian pasture soils.
- 97. Dimitrios Floudas Evolutionary aspects of atromentin synthesis genes in Agaricomycetes
- 98. Diana Navrátilová Spatial heterogeneity of decomposition and fungal community composition within single *Quercus petraea* leaves
- 99. Kevin Geyer A comparison of methods for measuring the efficiency of microbial metabolism
- 100. Ton Gorissen Uniformly <sup>13</sup>C-Labelled Biomass Tracers Advances in <sup>13</sup>C-Techniques tracing changes in soil microbial processes and populations
- 101. Petra Havlíčková Effects of plants on the structure, function and diversity of bacterial communities
- 102. Jussi Heinonsalo Evidences on the ability of mycorrhizal genus *Piloderma* to use organic nitrogen and deliver it to Scots pine
- 103. Vincent Herve Ecology and diversity of oxalotrophic bacteria an in silico analysis
- 104. Björn Hoppe Fungal functional diversity and enzyme activity patterns in decaying logs of 13 temperate tree species in an in situ decomposition experiment
- 105. Aicha Asma Houfani Enzyme activities of aerobic (hemi)cellulolytic bacteria isolated from Algerian soils and compost
- 106. Dominika Chmolowska Cellulose was decomposed faster in fallow soil than in meadow soil because of a guicker start of the process
- 107. Sarah Johnston Fungus-Bacteria Interactions in Decomposing Wood
- Grit Kabiersch Detection of organotin compounds and degradation by litterdecomposing fungi
- 109. Katharina Keiblinger Efficacy of biochar and compost on remediation of copper contamination in vineyard soils effects on soil microbiology
- 110. Harald Kellner Fungal research on an artificial deadwood decomposition experiment in the German Biodiversity Exploratories
- 111. Jaroslav Kukla The influence of traditional agriculture on soil organic matter in tropical ecosystems of Papua New Guinea

- Iuliia Kyiashchenko The effect of soil fertility on fungal communities, enzyme activities and soil carbon dynamics in unmanaged forests
- 113. Sabrina Leonhardt Fungal extracellular enzyme activity and biomass in coarse woody debris of 13 tree species in the early phase of decomposition.
- 114. Katya Litova Studies on biodegradation of naphthalene and anthracene by Aspergillus glaucus strain isolated from Antarctic soil
- 115. Ashish Malik Microbial communities' fungal to bacterial dominance alters carbon cycling in soil
- 116. Tomas Martin-Bertelsen Towards linking fungal genes to chemical spectra from soil organic matter using machine learning
- 117. Tijana Martinović Structure of microbial communities in the environmentally exposed construction wood samples of different species
- 118. Itamar Melo Isolation and Screening of Highly Cellulolytic *Trichoderma* spp. from the Amazon Rainforest
- 119. Sophie Mieszkin Effect of wood extractives on wood-degrading microorganisms and importance of the ecological niche
- 120. Karolin Müller Turnover of Microbial Carbon in the Detritusphere
- 121. Cesar Nicolas Cuevas Organic N decomposition by fungal community under fertilized spruce forest
- 122. Naoise Nunan Carbon dynamics in Amazonian podzols under climate change
- 123. Michiel Op De Beeck Soil organic matter degradation by ectomycorrhizal fungi
- 124. Tim Philpott Fungal decomposition of fine roots in response to variable retention silviculture
- 125. Sebastian Preusser Reciprocal Soil Transfer Experiments Improve the Understanding of Biological Regulation of Subsoil C-cycling
- 126. Salvador Rodríguez Zaragoza Recovering of soil protozoan trophic groups after a strong pulse of hydrocarbon contamination
- Mikhail Semenov DNA-based determination of soil microbial biomass carbon under conditions of restricted applicability of substrate-induced respiration and fumigationextraction
- 128. Sarker Mohammad Shakil Characterization of Fe<sup>3+</sup> reductants secreted by the closely related ectomycorrhizal fungus Paxillus involutus and the saprotrophic fungus *Hydnomerulius pinastri* during Fenton-based decomposition of organic matter
- 129. Ana Margarida Soares Bridging the priming effect into aquatic systems Primary producer-C stimulates the fungal decomposition of submerged litter
- 130. Florian Strasser Influences of carbon substrates and nitrogen availability on microbial-mediated cellulose degradation in an Austrian beech forest soil
- 131. Lucie Štercová Fungal biodiversity of wood decomposing species in national nature reservation of Salajka
- 132. Rodrigo Taketani Litter decomposition in mangroves the role of microbes revealed by DNA and mRNA sequencing
- 133. Aloysius Teo Using teabags to estimate decomposition rates across primary and secondary tropical forests, and investigating the functional role of termites
- 134. Vojtěch Tláskal Linking deadwood age with inhabiting bacterial community
- 135. Tomas Větrovský Sequence processing fast and easy SEED a GUI based user friendly sequence editor and pipeline for high-throughput amplicon processing

- 136. Alexandra Wolf Agro-ecosystem type and soil aggregate size impact soil carbon dynamics
- 137. Ilya Yevdokimov Microbial immobilization and incorporation into DNA of inorganic <sup>33</sup>P-labelled phosphorus
- 138. Lucia Žifčáková Correlation of lignocellulolytic genes expression and their activity in ME fungal cultures

### Soil Biogeochemistry and Nutrient Cycling

- 139. Roey Angel Optimizing the toolbox to investigate free-living diazotrophs in soil from bulk measurements to single-cell analysis.
- 140. Doreen Babin Effect of phenanthrene on the release of mobile organic matter and the bacterial community structure in soil
- 141. Paul Bodelier Unexpected stimulation of soil methane uptake by bio-based residue application An emerging property of agricultural soils offsetting greenhouse gas balance
- 142. Runa Boeddinghaus Land-use intensity and physico-chemical soil properties have distinct effects on microbial communities and enzyme activities of grassland soils
- 143. Ivano Brunner Shifts of C and N isotopes in fruiting bodies of fungi after 12 years of irrigation of a semi-arid pine forest
- 144. Elisa Catão Ammonia oxidizers in a non-nitrifying Brazilian savannah soil
- 145. Maria Cucu Different agricultural practices drive aerobic and anaerobic ammonia oxidisers niche segregation in a temperate paddy soil
- 146. Andreas Demey Impact of bioavailable phosphorus on plant and soil microbial communities in grassland under restoration management
- 147. Stefan Forstner How does long-term nitrogen input influence stoichiometric relationships between soil microbes and their resources?
- 148. Bo Fu The effect of temperature on the carbon isotope value of acetate in Philippine rice field soil
- 149. Ahlam Hamim Phosphate solubilizing microorganisms isolated from root and rhizosphere soil of ericaceous shrubs in the north of Morocco.
- 150. Christine Heuck Soil microbial biomass C:N:P stoichiometry and microbial use of organic phosphorus
- 151. Jiezhong Chen The Expression Analysis of Plasma Membrane Aquaporin Gene EiPIP2 in Eriobotrya japonica After AM Fungi Inoculation
- 152. María Irisarri Do soil type, rice cultivar and water management affect the bacterial denitrifying community of a paddy soil?
- 153. Sheku Kanu Interactive effects of Bacillus subtilis and seaweed (kelpak) on the growth, metabolites and yield of potato (Solanum tuberusom L.) under glasshouse conditions
- 154. Késia Lourenço Effected of regular or concentrate vinasse on greenhouse gases emissions from soil with sugarcane
- 155. Sven Marhan Nutrient limitation of soil microorganisms effects of grassland landuse intensity

- 156. Dora Neina Restoring the functional integrity of a Technosol with native organic materials
- 157. Maximilian Nepel Identifying potential key players of N2 fixation in European biological soil crusts
- 158. Laurent Philippot Recently identified microbial guild mediates soil N2O sink capacity
- 159. Frank Rasche Lasting influence of biochemically contrasting organic inputs on abundance and community structure of total and proteolytic bacteria in tropical soils
- 160. Karolina Tahovská Microbial activity in the context of acid deposition field manipulations with sulphur and/or nitrogen inputs to the forest soils
- 161. Irina Tanuwidjaja Influence of different clay minerals on the microbiome of soils and its functionality in simplified artificial systems
- 162. Cecile Thion Predicting temporal and spatial variations in bacterial phylogenetic and phenotypic community structure in glacier forefield chronosequences
- 163. Yang Zhou The functional profiles of soil microbial communities are determined by soil chemical properties but not community composition

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- 164. Pilar Andrés Effect of biochar application to soil on soil microbial communities structure and feeding habits a field study in Mediterranean soils
- 165. Olubukola Babalola Metal tolerant, plant growth promoting soil bacteria protected plants against the toxic effects of heavy metals (Cd, Cr, and Ni)
- 166. Guillaume BAY Effects of cropping system, depth, and sampling time on soil microbial communities
- 167. Andrew Bissett Effects of temporal pH shifts on ammonia oxidiser community structure and function
- 168. Ian Clark Response of Bacterial and Archaeal nitrifying populations to changing landscapes
- 169. Benjamin Costerousse Characterization of the bacterial processes responsible for zinc solubilization in wheat rhizosphere
- 170. Florine Degrune The influence of soil tillage on microbial communities changes along the soil profile
- 171. Anderson Ferreira Soil bacterial community under integrative production system at biomes savanna and Amazon
- 172. Davide Francioli Rhizosphere microbiome, plant community and soil nutrient availability a new approach to survey the bacterial assemblage in soil
- 173. Mercedes García Sánchez Digestate and fly ash applications in agricultural soils impact in the biomass and biodiversity of fungal communities.
- 174. Aurelia Gebala Does Land-Use Intensity Influence Microbial Resource Partitioning and Microbial Colonization Strategies of Organo-Mineral Complexes in Grassland Soils?
- 175. Mariangela Girlanda Plant genotype control over the recruitment of the tomato fungal microbiota
- 176. Daniel Graf Community assembly processes of N2O reducing prokaryotes in the rhizosphere- effect of edaphic factors and plant species

- 177. Yian Gu Pathogen-induced shifts in exudation alter the rhizosphere microbiome
- 178. Moritz Hallama Soil Microbial Phosphorus Dynamics are Affected by Cover Crops and Minimum Tillage
- 179. Penny Hirsch Abundance and activity of soil microbial communities revealed by metagenomics and metatranscriptomics
- 180. Anna-Sofia Hug Soil microbial diversity patterns at Sites of the Swiss Soil Monitoring Network
- 181. Christopher Jones Field-scale spatial variation in co-occurrence patterns of ammonia and nitrite oxidizing communities.
- 182. Milko Jorquera Exploring rhizobacterial community composition associated with plants grown in Chilean extreme environments using 16S rRNA-based molecular approaches
- 183. Hans-Martin Krause Influence of soil management history on microbial N2O production and reduction
- 184. Iva Krizkova-Kudlikova Characterization of Actinomycetes Antagonistic to Streptomyces spp.
- 185. Martin Krsek Bead-beating and isolation of environmental nucleic acids
- 186. Volery Lara Effects of management on soil microorganism communities in Swiss vineyards
- 187. Guillaume Lentendu Alfalfa root symbionts under soil nutrient pressure cooperation or competition?
- 188. Hongwei Liu Activation of salicylic acid defence signalling pathway reduced Archaea abundance and genes involved in nitrogen and carbon cycling in wheat rhizosphere
- 189. Pawel Lycus Newly isolated denitrifiers from low and high pH soil show little correlation between genotype and phenotype
- 190. Jarmila Makovníková The potential of agroecosystem services in relation to land use and biodiversity
- 191. Lokeshwaran Manoharan Enzymes related to organic matter degradation and agricultural management
- 192. Shinsuke Mori Changes in the oxidation-reduction potential and in bacterial profiles in the soil around direct-seeded rice under submerged conditions
- 193. Esther Muema Biochemically contrasting organic inputs combined with mineral nitrogen fertilizer shape the temporal variation of ammonia-oxidizing prokaryotic communities in an agricultural soil
- 194. Mary Musyoki Soil type, season and crop growth stage exert a stronger effect on rhizosphere microbial dynamics than the fungal biocontrol agent Fusarium oxysporum f.sp. strigae
- 195. Lidia Nicola Fumigation with Dazomet modifies soil bacterial and fungal communities in soil of apple orchards affected by Specific Replant Disease
- 196. Ansa Palojärvi Improved general plant pathogen suppressiveness by agricultural management practices
- Martina Putz Long-term nitrogen fertilization affects microbial communities regulating N2O emissions in arable soils
- 198. Vivian Rincon Florez Impact of strategic tillage on nitrogen cycle genes (amoA and nifH) in no-till systems in Queensland, Australia
- 199. Jenna Ross Isolation of a novel ammonia oxidising archaeon, representative of the Nitrososphaera 'sister' lineage

- 200. Suikinai Nobre Santos Annotation of gene cluster Involved phenazine biosynthesis in Streptomyces CMAA 1322 also too structural elucidation of 1,6 dimethoxyphenazine.
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- 202. Alise Senberga Evaluation of Effectiveness of Rhizobia and Plant Interaction in Different Soil Types
- 203. Abhi Shah Microbial activity along a continuous subsurface core from an agriculture field at the estuarine region of Mahi river: correlation with sediment characteristics
- 204. Hannes Schmidt Diversity and spatial distribution of diazotrophs associated with micro-environments of wetland rice
- 205. Elvira Schnyder Methanotroph diversity increases methane oxidation
- 206. Susanne Schreiter Roots of decline? The SARISA project
- 207. Magdalena Steiner Microbial diversity and ecosystem functioning in vineyards
- 208. Nicolas Theodorakopoulos Microorganism's enzymes implication in nitrous oxide emissions in a natural agricultural field at a fine time scale study
- 209. Maaike van Agtmaal Exploring the phytopathogenic seedbank of agricultural soils diversity of soil borne plant pathogens in relation to edaphic properties and the soil microbial community
- 210. Wu Xiong Effects of black pepper-vanilla rotation on vanilla rhizosphere fungal communities in relation to Fusarium wilt disease
- 211. Tianjie Yang Trophic network architecture of root-associated bacterial communities determines pathogen invasion and plant health
- 212. Qing Yao The functional profiles of soil microbial communities are determined by soil chemical properties but not community composition
- 213. Judith Zimmermann The biocontrol agent Fusarium oxysporum f.sp. strigae its detection and effects on beneficial indigenous microorganisms in a maize rhizosphere
- 214. Josephine Zimudzi The microbial communities associated with potato rhizosphere under different seasonal conditions in South Africa

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- 215. Sarah Addison Soil microbes and their importance in shaping our forests using gPCR
- 216. Toke Bang-Andreasen Responses in Active Microbial Communities and Expression of Important Functional Genes in Forest and Agricultural Field Soil after Wood Ash Addition Revealed by Metatranscriptomics
- 217. Tommaso Bardelli Complex effects of altitude and exposure on microbial communities in (sub)alpine soils
- 218. Felipe Bastida Landscape proteogenomics explaining the functional-phylogenetic relationships of microbial communities by gradients of organic C availability in soil
- 219. Garazi Benito Carnero Tree species effect on soil microbial community
- 220. Monique Carnol Carbon substrate utilization and microbial biomass in European forest soils are related to tree species diversity
- 221. Carles Castaño Soler Drying treatment of soil samples affects DNA recovery but does not change the fungal community structure by metagenomic analysis

- 222. Carla Cruz-Paredes Using fungal and bacterial growth to evaluate the effects of ash application on forest soils
- 223. Timo Domisch Winter in a changing climate affecting the survival of Scots pine seedlings
- 224. Beat Frey Pyrosequencing based assessment of bacterial and fungal community compositions in compacted and regenerated forest soils
- 225. Anna Frymark-Szymkowiak Soil β-Glucosidase activity under canopy of White Poplar in riparian forests.
- 226. Kezia Goldmann Spatial variation of the fungal metagenome in temperate beech forests across Germany
- 227. Erika Gömöryová Soil microbial community changes in the disturbed Norway spruce stands during a 10-years period
- 228. Sue Grayston CH4 and N2O microbial communities respond to site preparation and fertilization in wet forests
- 229. Michal Choma Recovery of ectomycorrhizal community of a boreal forest after three decades of N fertilisation
- 230. Leticia Izquierdo A new promising molecular marker to study the functional diversity of fungal communities the glycosyl hydrolase 63 gene
- 231. Veronika Jílková Methane flux in wood ant (Formica polyctena) nests and the surrounding forest floor
- 232. Jan Kopecký Metabolite profiles of soil actinobacteria follow their phylogeny and environmental factors at the isolation sites
- 233. Diego Leiva Cáceres Relations between Peltigera lichen's derived factors and its associated bacterial communities
- 234. Sandrine Malchair Spatial variability of soil microbial processes in a temperate mixed forest
- 235. Oscar Martínez Fungal communities associated with rhizosphere of Nothofagus alpina from different volcanic ash-derived soils in southern Chile
- 236. David Myrold Development and Decline of Microbial Communities Associated with Ectomycorrhizal Mats
- 237. Pascal Nassal The importance of fungal-fungal and bacterial-fungal interactions for phosphorus dynamics in forest soils
- 238. Mike Ogden Fine scale modification of soil physical properties by fungi reinforcement and repellency in the hyphosphere
- 239. Jade O'Leary Multi-dimensional mycelia interactions
- 240. Xavier Parlade Dynamics of Boletus edulis extraradical soil mycelium and sporocarp production in managed forests
- 241. Taina Pennanen Virus host switches between pathogenic, mycorrhizal and saprotrophic fungal species in a boreal forest
- 242. Camilla Pereira Arbuscular mycorrhizal fungi in protected areas of northeastern Brazil
- 243. Diogo Pinho A first look at the Quercus suber (cork oak) root microbiome differences between healthy and declined trees
- 244. Flavia Pinzari Overlap in the metabolic functions of cellulose-decomposing leaf litter fungi
- 245. Tereza Poláčková Ecology of soil yeast communities in mixed temperate forest soils

- 246. Daria Rapoport Isolation and cultivation of actinobacteria from acid soils with high occurrence of Trebon clade
- 247. Ana Rincón Fire recurrence effects over the structure and activity of ectomycorrhizal fungal communities in Mediterranean pine forests
- 248. Alice Roy-Bolduc High richness of ectomycorrhizal fungi and low host specificity in a coastal sand dune ecosystem
- 249. Minna Santalahti Revealing sources of biological methane production in boreal upland forest
- 250. Outi-Maaria Sietiö The effect of photosynthesis-derived C flow on the microbial community structure and enzymatic activities in boreal forest
- 251. José A. Siles Archaeal, bacterial and fungal abundance and diversity along an altitudinal gradient in Alpine forest soils
- 252. Izabela Sondej Impact of wild boar (Sus scrofa) rooting on the soil seed bank in Białowieża Forest.
- 253. Martina Štursová Spatial heterogeneity of mountainous soil is associated with high beta diversity of microbial community
- 254. Martina Vašutová Mycorrhizal community structure across an alpine tree line ecotone
- 255. Christina Weißbecker Patterns of soil fungal communities in subtropical Chinese forests in relation to plant diversity

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- 256. Ina Alsina The yield of onions and its quality depending on mycorrhiza inoculation
- 257. Lucas Braga Earthworm-microbe interaction can be associated to less harsh conditions in green sugarcane systems
- 258. Ana Correia Carbon source and availability influence the production of antimicrobial compounds
- 259. Alper Dede Rhizosphere of olive tree: a source of plant growth promoting bacteria
- 260. Laila Dubova The effects of michorrhyza fungy on the tomatoe plant water retention ability
- 261. Iva Cholakova Characterization of plant-associated bacteria isolated from highly drought tolerant Pistacia therebinthus
- 262. Polina Ivanova Antibiotic activity of actinobacteria associated with millipedes and earthworms
- 263. Xianwang Kong Effects of the nitrification inhibitor 3,4-dimethylpyrazole phosphate (DMPP) on N2O emissions from clover residues and interaction with the earthworm Lumbricus terrestris
- 264. Tat'yana Kotova Suppressive activity of the intestinal fluid of diplopods against yeasts
- 265. Ines Mandic Mulec Kin discrimination between sympatric soil isolates of Bacillus subtilis
- 266. Lucas Mendes The role of rhizosphere microbiome in soilborne fungal disease suppression in common bean
- 267. Anna Rawlings Decay in the canopy
- 268. Ruth Schmidt Microbial airborne talk effect of fungal volatiles on bacteria

- 269. Ana Soares Specificity and biotic local selection in Streptomyces interactions influences antimicrobial activity
- 270. Tamara Těšitelová Two widespread green Neottia species (Orchidaceae) show mycorrhizal preference for Sebacinales in various habitats and ontogenetic stages
- 271. Sannakajsa Velmala Function by form a tentative insight to the link between growth and the diversity of ectomycorrhizal fungi
- 272. Zhihui Xu Comparative proteomics analysis of Bacillus amyloliquefaciens SQR9 revealed the key proteins involved in in situ root colonization

# **Microbes in the Changing Environment**

- 273. Abdulmajeed Al Khajeh Fish emulsion as a food base for halophilic actinomycetes promoting growth of Salicornia bigelovii in a sandy soil in the United Arab Emirates
- 274. Rana Shahbaz Ali Effects of substrate complexity and temperature on growth of different microbial groups
- 275. Sawa ARAI Carbon-starvation in light induced tolerance to hyperosmotic stress in purple photosynthetic bacterium Rhodopseudomonas palustris
- 276. Andrea Borsodi Microbial communities inhabiting the rhizosphere of halophyton plants living nearby Hungarian soda lakes
- 277. Tabitha Bucher Disturbance of the bacterial cell wall specifically interferes with biofilm formation
- 278. Cristina Cruz Alleviating the N limitation expands the possibilities for structuring soil bacterial communities - evidence based on the impacts of 5 years' manipulation of N dose and form in a Mediterranean ecosystem
- 279. Jonathan De Long Soil microbial and nematode communities respond differently to warming and plant functional group removal across a post-fire boreal forest successional gradient
- 280. Teresa Dias Impacts of N enrichment on Mediterranean biological soil crusts community and functions the unseen evidence from soil pigments
- 281. Francisco Dini-Andreote Unveiling the blueprint of marine-terrestrial transition in bacterial adaptation and evolution
- 282. Dana Elhottová Bacterial antibiotic resistance heterogeneity in natural subterranean habitats
- 283. Huyuan Feng Influences of Long-term Nitrogen Fertilization on Fungal Endophyte Community of Three Grasses in an Alpine Meadow
- 284. Nawras Ghanem The transport of marine phages in soil as a tool of understanding the interaction of surface-subsurface events
- 285. Osnat Gillor Hydration dynamics in desert soil mediate antagonism of actinobacteria
- 286. Sydney Glassman Ectomycorrhizal fungal spore bank recovery after a severe forest fire: Some like it hot
- 287. Catarina Gouveia Effect of Increased N availability on Ammonium oxidizing bacteria populations A possible Bioindicator in Mediterranean ecosystem
- 288. Kelly Gravuer Phylogenetic estimation of ecologically important traits illuminates microbial community responses to change in natural and agro-ecosystems

- 289. Stefan Green Methodological Improvements to Amplicon-Based Surveys of Microbial Community Structure
- 290. Jeremiah Henning Fungi on mountainsides contrasting elevational and seasonal patterns among root-associated fungal groups.
- 291. Andreas Herrmann Spatial and seasonal variability of the microbial community in forest fen soils on North-East-Germany
- 292. Jana Judova The Contingency of some Biotic and Abiotic Parameters in Arable Land and Permanent Grasslands
- 293. Tatiana Khomutova Characterization of microbial pool in sub-kurgan paleosols of different ages in desert-steppe zone in relation to the holocene dynamics of climate
- 294. Sara König Hot spots and cold spots modelling biodegradation dynamics under disturbance regimes
- 295. Ramóna Kovács Characterization of mycorrhizophere in a Hungarian saline-sodic grassland
- 296. Anna Kuznetsova The first study of actinomycetes complexes in Prietonie region soil
- 297. Xiaofei Lyu Soil bacterial community along a successional series of tidal flats in the Yellow River Delta
- 298. Aurora MacRae-Crerar Ecological determinants of soil bacterial community structure across multiple scales in a Mongolian global change experiment
- 299. Marta Misiak Soil fungal responses to warming in polar regions
- 300. Luis Morgado Compositional shifts in arctic ectomycorrhizal fungal community in response to long-term increased snow depth in Northern Alaska
- 301. Eric Morrison Soil warming changes litter chemistry and fungal community composition but not decomposition rate
- 302. Sally Otto Catch me if you can The impact of mycelia-based dispersal on predator-prey interactions and biodegradation of soil contaminants
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- 304. Elizaveta Pershina Looking for the core microbiome of the main types of soils in Russia
- 305. Kristin Rath The influence of salinity on saprotrophic fundi and bacteria in soil
- 306. Klára Řeháková Potential activity of microbial community in the Biological Soil Crusts
- 307. Nermina Saronjic How do soil microbial communities react on droughts and heavy rainfall events?
- 308. Philipp-André Schmidt Response of Soil Fungal Communities to Extended Drought
- 309. Frank Solano-Campos Preliminary data of soil nematode communities along a rainfall gradient in Costa Rica.
- 310. Afnan Suleiman Active microbial community resilience in disturbed soil with nitrogen source enrichment and nitrification inhibitor
- 311. Tibor Szili-Kovács Genetic diversity and catabolic activity profiles of rhizosphere bacterial communities during dry and wet seasons in a solonchak grassland, Hungary
- 312. Alexandra Šimonovičová Spatial distribution of microscopic fungi under old environmental burdens
- 313. Adam Šťovíček Bacterial response to rainfall and draught cycles in desert soil
- 314. Daniela Trojan Investigating the ecophysiology of the ubiquitous Acidobacteria in the dynamic soil environment
- 315. Tushar Yadav Effect of Cosmetic Based Nanowaste on Sludge and Soil Microflora

- 316. Baogui Zhang Response characteristics of soil microorganism to permafrost degradation in the upstream regions of the Shule river basin, Qinghai-Tibetan Plateau
- 317. Junling Zhang Unexplored Biodiversity and Function of arbscular mycorrhizal fungion the Tibetan Plateau

### Microbial Life in Contaminated and Anthropogenic Soils

- 318. Valeria Ancona Microbiological indicators to evaluate soil quality of degraded areas in Southern Italy after compost addition
- 319. Thomas Banitz Spatial metrics indicate bacterial degradation benefits from mycelial networks
- 320. Angelantonio Calabrese Use of molecular techniques to characterize the microbial communities for soil ecology assessment in degraded sites.
- 321. Juan Campos Soil dehydrogenase activity under the presence of some exobiotics. A toxicity index is proposed.
- 322. Stefano Covino A pyrosequencing-based metagenomic study of microbial communities during co-composting of creosote-impregnated wood and green wastes
- 323. Sabrina Festa Monitoring the impact of bioaugmentation with a PAH-degrading strain on different soil microbiomes using pyrosequencing
- 324. Alena Filipová A novel bioaugmentation approach for PAH-degrading bacteria in soil: Adaptability as assessed by molecular biology techniques
- 325. Fanny Flores Microbial activity of chromium polluted soil from Guanajuato México, during in situ biostimulation assay.
- 326. Ulrike Gerber Interactions of natural occurring eukaryotic microorganisms with uranium(VI)
- 327. Paola Grenni Effects of compost addition and Medicago sativa occurrence on PCB biodegradation in a historically contaminated soil
- 328. Anna Grobelak The possible application of microorganisms in promoting plant growth and improving plant biomass in the phytoremediation of anthropogenic and contaminated soils
- 329. María Gutiérrez Núñez Non-target effects of pesticides on the microbial activity in agricultural soil
- 330. Lenka Harantová Factors influencing microbial community development during primary succession on spoil heaps after brown coal mining
- Karin Hellauer Sequential managed aquifer recharge leads to a high diverse microbial community resulting in a better attenuation of moderate degradable trace organic chemicals (TOrC)
- 332. Anne Houles Ecological restoration of nickel mine sites in New Caledonia Characterisation of ectomycorrhizal fungal community the key enabling the monitoring of facilitation process between plants.
- 333. Alica Chroňáková Boreal acid sulphate soils changes in bacterial communities along vertical profile and between total and active pools
- 334. Inge Jambon Bioremediation of chlorendic acid, a highly chlorinated organic pollutant, by exploiting a fungal-bacteria consortium native to the contaminated field

- 335. Michal Kaminski Omics approach in analysis of Pseudomonas mandelii ssp. capable of bioaccumulating hexachlorocyclohexane
- 336. Hülya Kaplan Diversity of bacteria involved in 13C-labelled wheat root decomposition and efflux-mediated metal resistance in metal contaminated soils remediated with amendments
- 337. Pawel Krawczyk Bioinformatic approach to analysis of plasmid pool in metagenomes from polluted soils
- 338. Jennifer Mesa Marin Ecology of soil bacteria in bioremediation indigenous plant growth promoting rhizobacteria in native Spartina maritima as a tool for the restoration of heavy metal polluted salt marshes
- 339. Zuzana Michalkova Interactions of nano zerovalent iron with Acidithiobacillus ferooxidans Implications for soil remediation
- 340. Annett Mikolasch Oil-degrading bacteria isolated from the rhizosphere of plants growing in oil-contaminated soils from Kazakhstan
- 341. Irma Morelli Pyrosequencing reveals bioaugmentation impact on the dynamics of bacterial community on phenanthrene-contaminated soil
- 342. Marta Moreno Valencia Effect of olive and vine wood ashes on the dehydrogenase activity in a crop land.
- 343. Martina Plačková Bacterial community characteristics under decades-lasting antibiotics selection pressures
- 344. Thomas Pommier Response of soil microbial community to titanium dioxide nanoparticles a cascading pitch on the nitrogen cycle
- 345. Pavla Průchová Composition and activity of microbial communities in soil contaminated by heavy metals
- 346. Rashmi Saikia Influence of sources of carbon in growth media on the yield of biosurfactant by the microbe isolated from crude oil contaminated soil
- 347. Hokyung Song Abandoned tropical tin mine site shows chnges in microbial community with restoration
- 348. Gangavarapu Subrahmanyam Abundance and diversity of ammonia oxidizing archaea and bacteria in long-term industrial effluent polluted soils, Gujarat, Western India
- 349. Karel Švec Composition of fungal and bacterial communities in mercury polluted areas
- 350. María Touceda-González Molecular characterization of the rhizobacterial communities of two Ni-hiperaccumulating subspecies of Alyssum serpyllifolium endemic to the Iberian Peninsula.
- 351. Gustavo Valdecantos Biotic and abiotic factors affect the colonization and the dynamics of bacterial community assemblage in irradiated soil microcosms
- 352. María Vásquez Murrieta Metal tolerance and biosorption potential of endophytic fungi isolated from Bahia absinthifolia
- 353. Carl-Eric Wegner Persisting in slag insights into aluminium resistance from early industrial mineral leaching
- 354. Franco Widmer Effects of different nanoparticles on soil microbial community structures and plant-microbe interactions
- 355. Yucheng Wu Responses of Thaumarchaeotal community in agricultural soils to acidification and polycyclic aromatic hydrocarbons contamination

356. Lijuan Yan - Comparative phylogenetic analysis of bacterial community dynamics during multi-year bioremediation of oil-contaminated soil in a boreal climate

# Archaeomicrobiology, Paleomicrobiology and Microbial Forensics

- 357. Cecile Gubry-Rangin Molecular adaptation of the ammonia monooxygenase amoA gene during the ancient and rapid diversification of terrestrial Thaumarchaeota
- 358. Eline van Asperen Establishing dung fungal spores as a proxy for herbivore abundance an experimental approach