

AperTO - Archivio Istituzionale Open Access dell'Università di Torino

PERN: an EU-Russia initiative for rhizosphere microbial resources

This is the author's manuscript

Original Citation:

Availability:

This version is available <http://hdl.handle.net/2318/1520996> since 2017-05-11T18:39:46Z

Published version:

DOI:10.1016/j.tibtech.2015.03.005

Terms of use:

Open Access

Anyone can freely access the full text of works made available as "Open Access". Works made available under a Creative Commons license can be used according to the terms and conditions of said license. Use of all other works requires consent of the right holder (author or publisher) if not exempted from copyright protection by the applicable law.

(Article begins on next page)

This Accepted Author Manuscript (AAM) is copyrighted and published by Elsevier. It is posted here by agreement between Elsevier and the University of Turin. Changes resulting from the publishing process - such as editing, corrections, structural formatting, and other quality control mechanisms - may not be reflected in this version of the text. The definitive version of the text was subsequently published in TRENDS IN BIOTECHNOLOGY, 33 (7), 2015, 10.1016/j.tibtech.2015.03.005.

You may download, copy and otherwise use the AAM for non-commercial purposes provided that your license is limited by the following restrictions:

- (1) You may use this AAM for non-commercial purposes only under the terms of the CC-BY-NC-ND license.
- (2) The integrity of the work and identification of the author, copyright owner, and publisher must be preserved in any copy.
- (3) You must attribute this AAM in the following format: Creative Commons BY-NC-ND license (<http://creativecommons.org/licenses/by-nc-nd/4.0/deed.en>), 10.1016/j.tibtech.2015.03.005

The publisher's version is available at:

<http://linkinghub.elsevier.com/retrieve/pii/S0167779915000670>

When citing, please refer to the published version.

Link to this full text:

<http://hdl.handle.net/2318/1520996>

PERN: an EU–Russia initiative for rhizosphere microbial resources

- [Stéphane Declerck¹](#),
- [Anne Willems²](#),
- [Marcel G.A. van der Heijden^{3, 4, 5}](#),
- [Giovanna Cristina Varese⁶](#),
- [Olga Turkovskaya⁷](#),
- [Lyudmila Evtushenko⁸](#),
- [Irena Ivshina^{9, 10}](#),
- [Philippe Desmeth¹¹](#)

Show more

[doi:10.1016/j.tibtech.2015.03.005](https://doi.org/10.1016/j.tibtech.2015.03.005)

Millions of microbial taxa inhabit the rhizosphere and could be used as biofertilizers, biopesticides, and/or for bioremediation. Only a fraction of these microbes have been described and/or are being utilized. Most are dispersed in collections, but coordination of their accessibility and availability is challenging. Here, we present the Pan-European Rhizosphere Resource Network (PERN), which is a transnational repository of microorganisms whose objectives are to facilitate access to rhizosphere resources and information and help users with technical and legal issues.

Keywords

- access and benefit sharing;
 - biopesticide;
 - biofertilizer;
 - bioremediation;
 - culture collection;
 - rhizosphere;
 - Nagoya protocol;
 - plant microbiome
-

The soil microbial reservoir and the rhizosphere microbiome

Soil contains a tremendous diversity and abundance of microbes, such as phages and viruses (up to 10^9 g⁻¹ soil), bacteria ($4\text{--}20 \times 10^9$ cells cm⁻³), protists ($10^4\text{--}10^7$ cells m⁻²), and fungi ($6.9 \times 10^6\text{--}2.1 \times 10^9$ cells g⁻¹) with millions of species or ecotypes from each group [1](#), [2](#) and [3](#). The abundance and interactions of microbes largely depend on the environment, and microbes have a big impact on ecosystems, principally determining agroecological functioning and stability [\[4\]](#). Many of these soil microbes inhabit the rhizosphere, that is, ‘the interface between roots and soil where interactions among microbes and invertebrates affect biogeochemical cycling, plant growth and tolerance to biotic and abiotic stress’ [\[5\]](#).

The need to preserve rhizosphere microbial diversity in culture collections

There are various reasons to preserve rhizosphere microbial diversity in a culture collection. A range of root-associated microbes has been reported to be beneficial for plant growth and nutrition [6]. For instance, estimates suggest that mycorrhizal fungi are responsible for up to 75% of the phosphorus that is acquired by plants annually, and many legumes rely on associations with nitrogen-fixing bacteria for nitrogen acquisition [6]. Moreover, root-associated microbes can improve plant resistance to pests and diseases [7]. This emphasizes the need to collect and preserve root-associated microbes because they can be used as biofertilizers, biopesticides, and/or for bioremediation.

The recent application of large-scale environmental DNA sequencing has revealed that only a fraction of microbes in soil or in the rhizosphere has been cultured by traditional techniques and a significant number of abundant microbial groups, such as the *Acidobacteria*, *Planctomycetes*, and *Verrucomicrobia*, are underrepresented among cultivated strains [8]. However, as more varied cultivation conditions are tested, representatives of these previously uncultured taxa are gradually being cultured [9]. These efforts should be continued, scaled up [10], and coordinated. A promising development in this respect is the recent success of culturomics (a method using high-throughput culture to enable the description of the microbial composition [11]) in human gut microbiome studies, which demonstrated a greater culturability when diverse culturing conditions were applied [12].

It is important that the microorganisms, which have been isolated at considerable expense, be preserved in optimal conditions so that they are available for future research and for the development of applications. This is not a trivial issue because many environmental strains may be sensitive to current preservation techniques [13]. The expected increase in the number of species and strains available in culture will require the simultaneous development and improvement of preservation methods to guarantee viability and optimal preservation of sensitive microorganisms without a change in their morphology, physiology, and genetics until use [14].

No single laboratory is currently set up to isolate, identify, preserve, characterize, or have an extensive knowledge of, all rhizosphere microorganisms. Microbial isolates may disappear from individual laboratories once research funding has ceased, emphasizing the need to preserve isolates in a coordinated effort and to promote exchange and cooperation between laboratories.

The Pan-European Rhizosphere Resource Network

In 2010, the European FP7 project BRIO was initiated to set up a virtual network of rhizosphere microbiological resource centers from Western Europe and Russia. The objective was to organize cooperation between specialized collections containing beneficial microorganisms isolated from the rhizosphere and to constitute a virtual, common, wide-range pool of microbial rhizosphere diversity that was exploitable for research and industry: PERN (<http://www.PERN-BRIO.eu>). PERN will ease the access to rhizosphere resources and information located in different collections via a single portal. It will also help users to comply with legal issues (Box 1) arising from: (i) the Convention on Biological Diversity (CBD; <http://www.cbd.int>); (ii) the Nagoya Protocol (NP) on Access and Benefit Sharing (ABS; <http://www.cbd.int/abs/>); and (iii) national and regional laws implementing this Protocol, such as the European Union (EU) Regulation 511/2014.

Box 1.

Access and benefit sharing

ABS is a concept coined in the CBD and ruled by the NP under the CBD. Access to biological resources (e.g., deposited in culture collections) is granted provided that a fair and equitable share of generated benefit returns to the party providing these resources. Regulation EU 511/2014 translates the NP into European law.

Nagoya Protocol

The NP is a protocol that was adopted on October 29, 2010 and entered into law on October 12, 2014. It provides a legal framework for the fair and equitable sharing of benefits arising from the utilization of genetic resources, thus contributing to the conservation and sustainable use of biodiversity.

Material accession agreement

An MAA is a contract between the depositor of a microbial sample and the collection where the sample is preserved. This document records minimal data required by law and necessary for scientific labeling, such as place and date of sampling, etc. It specifies the respective rights and duties of the depositor and collection.

Material transfer agreement

An MTA is a contract between the providing collection and the purchaser of biological samples. 'MTA' is a generic term that includes shipment document, standard delivery notice, standard invoice containing standard requirements, or more specific contracts, including tailor-made mutually agreed terms.

Definitions available from <http://bccm.belspo.be/projects/mosaicc> and <http://bccm.belspo.be/projects/trust>

PERN is organized as a virtual collection, centrally managed in Brussels by the Belgian Coordinated Collections of Microorganisms (BCCM; <http://bccm.belspo.be/>) and enriched with microbes provided by culture collections in Belgium [BCCM/MUCL (<http://bccm.belspo.be/about-us/bccm-mucl>) and BCCM/LMG (<http://bccm.belspo.be/about-us/bccm-lmg>)], Italy (Mycotheca Universitatis Taurinensis (MUT; <http://www.mut.unito.it/en>), Switzerland (The Swiss Collection of Arbuscular Mycorrhizal Fungi; <http://www.agroscope.admin.ch/bodenoekologie/08050/08067/08068/index.html?lang=en>), and a consortium of Russian collections [All-Russian Collection of Microorganisms (VKM; <http://vkm.ru/>); Institute of Ecology and Genetics of Microorganisms (IEGM; <http://www.iegm.ru/iegmcol/>); and the Institute of Biochemistry and Physiology of Plants and Microorganisms (IBPPM; <http://ibppm.ru/>)].

Each culture collection within PERN is responsible for its own microbial resources [identification; preservation; viability, identity, and purity (VIP) check; distribution; and legal issues; i.e.,

compliance with CBD and the NP) and the implementation of a data set that comprises a taxon and strain property browser (Figure 1). The central management in Brussels hosts and updates the PERN website, facilitates access to microbial resources via the unique PERN portal, and coordinates and guarantees common standards.

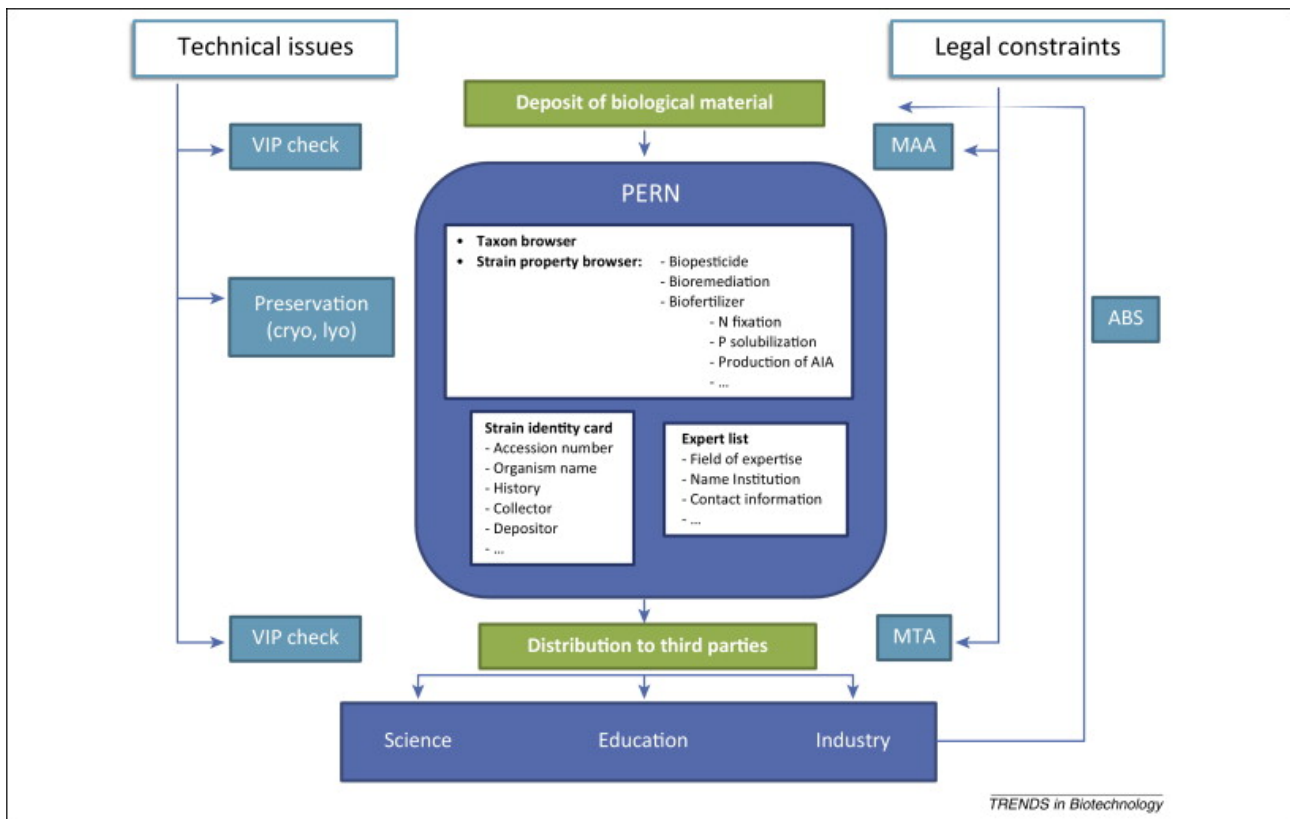


Figure 1.

Technical issues and legal constraints in the Pan-European Rhizosphere Network (PERN). Abbreviations: ABS, access and benefit sharing; AIA, acetic indole acid; MAA, material accession agreement; MTA, material transfer agreement; VIP, viability, identity, and purity.

Figure options

The total diversity within these collections currently exceeds 63 000 strains (many collected from food, feed, and agroenvironments), of which 1100 are rhizosphere microorganisms with documented properties available within PERN in compliance with international rules. These strains have been selected according to strict criteria for their biofertilizer potential (770 strains), biopesticide potential (81 strains), and bioremediation potential (291 strains). Their number is increasing constantly.

At strain deposit and distribution, a VIP check is performed (Figure 1). After the deposit of biological material, the organism is preserved following standard preservations techniques that differ from organism to organism, (i.e., by cryopreservation, lyophilization, or both). In some cases, storage under oil and water is also applied. Searches within this virtual collection are facilitated by a 'taxon browser' (e.g., accession number, organism name, etc.) and a 'strain property browser' (e.g., biofertilizer, biopesticide, and bioremediation). In both cases, an identity card presenting an exhaustive data set of information (Figure 1) is provided.

The website also presents a panel of international experts in applied microbial biotechnologies, such as biofertilizers, biopesticides, or bioremediation. An extensive list of protocols on how to isolate, culture, and preserve microbes from the rhizosphere as well as a list of documents (e.g., CBD, NP, EU as well as Russian legislation on intellectual Property Rights, and biosafety) is also provided to assist scientists and industries in the management and application of microorganisms.

The PERN collection welcomes other international collections that agree to the code of conduct set to maintain the quality of the material offered. The list of experts on rhizosphere microbiota will also be enlarged to include a wider range of expertise and cover more applications of rhizosphere microbes.

Access and use of microbial resources in PERN: legal constraints

For many stakeholders, access to microbial material is not only a technical issue, but also includes administrative and legal considerations: how to get the material through the due process and respecting the rights, including intellectual property rights, of third parties on the material [15]. Curators within microbial collections are increasingly asked to provide legal advice on possible legitimate uses and how to minimize the risk of lawsuits arising from the indiscriminate use of microbes. Regulations concerning biosafety and biosecurity are of primary importance for safeguarding public health. As major contributors to research and innovation (R&I), collections endeavor to deliver biological material with respect to intellectual property rights and benefit sharing. With the enforcing of the NP, collections are adapting their general conditions of distribution and deposit. To meet the new rules, more stringent data management, strict administrative procedures, quality management, and codes of conduct are required to guarantee the lawfulness of operations for the preservation, documentation, and distribution of microorganisms by the collections. Therefore, within PERN, a common policy for the deposition and distribution of microbial material and data has been set up in compliance with international regulations. This common policy is translated into standard documents for accession (deposition) of material into a member of the collection network (Material Accession Agreement, MAA) as well as for transfer (distribution) to third parties (Material Transfer Agreement, MTA) which represent a legally binding contract. This ensures that microbial resources acquired from a PERN member are not only technically effective, but also legally 'fit for use', providing the user with the legal certainty that they may commercialize the outputs of R&I without legal impediment (Box 1). PERN also proposes a selection of relevant legal texts from the EU and from the Russian Federation with translations and summaries as a service to microbiologists.

Concluding remarks and future perspectives

In the face of increasing numbers of microbes isolated from soil and rhizosphere and the development of methods to culture the 'unculturables', as well as more stringent international legislation, there is an urgent need for a coordinated international effort to preserve and distribute well-identified and characterized microorganisms that are legally fit for use in biotechnology. PERN was created to address these objectives by federating microbial culture collections in Western Europe and Russia hosting rhizosphere microbes. This virtual collection follows international legislation and is supported by a panel of scientific experts in the domain of rhizosphere microbiology and applications. The overall aim of PERN is to help manage rhizosphere microorganisms for sustainable agriculture and food production, and for developing diverse microbial environmental biotechnologies.

Acknowledgments

The 'Banking Rhizosphere Micro-Organisms European – Russian Initiative to set up a network of rhizosphere microbiological resources centres' (BRIO) was supported by the Seventh Framework Program FP7-KBBE-2010-4 of the European Commission under grant agreement No 266106. The authors are grateful to Kyle Hartman who checked the manuscript for English language use.

References

- [1](#) N. Fierer, *et al.* Comparative metagenomic, phylogenetic and physiological analyses of soil microbial communities across nitrogen gradients *ISME J.*, 6 (2012), pp. 1007–1017
- [2](#) R.D. Bardgett, W.H. van der Putten Belowground biodiversity and ecosystem functioning *Nature*, 515 (2014), pp. 505–511
- [3](#) M. Blackwell The Fungi: 1, 2, 3...5.1 million species? *Am. J. Bot.*, 98 (2011), pp. 426–438
- [4](#) M. Saleem, L.A. Moe Multitrophic microbial interactions for eco- and agro-biotechnological processes: theory and practice *Trends Biotechnol.*, 32 (2014), pp. 529–537
- [5](#) L. Phillipot, *et al.* Going back to the roots: the microbial ecology of the rhizosphere *Nat. Rev. Microbiol.*, 11 (2013), pp. 789–799
- [6](#) M.G.A. van der Heijden, *et al.* The unseen majority: soil microbes as drivers of plant diversity and productivity in terrestrial ecosystems *Ecol. Lett.*, 11 (2008), pp. 296–310
- [7](#) R.J. Rodriguez, *et al.* Fungal endophytes: diversity and functional roles *New Phytol.*, 182 (2009), pp. 314–330
- [8](#) U.N. da Rocha, *et al.* Exploration of hitherto-uncultured bacteria from the rhizosphere *FEMS Microbiol. Ecol.*, 69 (2009), pp. 313–328
- [9](#) U.N. da Rocha, *et al.* Cultivation of hitherto-uncultured bacteria belonging to the *Verrucomicrobia* subdivision 1 from potato (*Solanum tuberosum* L.) rhizosphere *J. Soil Sediment*, 10 (2010), pp. 326–339
- [10](#) V.H. Pham, J. Kim Cultivation of unculturable soil bacteria *Trends Biotechnol.*, 30 (2012), pp. 475–484
- [11](#) G. Greub *Culturomics: a new approach to study the human microbiome* *Clin. Microbiol. Infect.*, 18 (2012), pp. 1157–1159
- [12](#) J.C. Lagier, *et al.* Microbial culturomics: paradigm shift in the human gut microbiome study *Clin. Microbiol. Infect.*, 18 (2012), pp. 1185–1193
- [13](#) B. Vekeman, *et al.* A generally applicable cryopreservation method for nitrite-oxidizing bacteria *Syst. Appl. Microbiol.*, 36 (2013), pp. 579–584
- [14](#) K. Heylen, *et al.* Safeguarding bacterial resources promotes biotechnological innovation *Appl. Microbiol. Biotechnol.*, 94 (2012), pp. 565–574

[15](#) K. McCluskey Biological resource centers provide data and characterized living material for industrial biotechnology *Ind. Biotechnol.*, 9 (2013), pp. 117–122