



Towards a global platform for linking soil biodiversity data

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Title: Towards a global platform for linking soil biodiversity data

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Abstract

Soil biodiversity is immense, with an estimated 10-100 million organisms belonging to over 5000 taxa in a handful of soil. In spite of the importance of soil biodiversity for ecosystem functions and services, information on soil species, from taxonomy to biogeographical patterns, is incomplete and there is no infrastructure to connect pre-existing or future data. Here, we propose a global platform to allow for greater access to soil biodiversity information by linking databases and repositories through a single open portal. The proposed platform would for the first time, link data on soil organisms from different global sites and biomes, and will be inclusive of all data types, from molecular sequences to morphology measurements and other supporting information. Access to soil biodiversity species records and information will be instrumental to progressing scientific research and education. Further, as demonstrated by previous biodiversity synthesis efforts, data availability is key for adapting to, and creating mitigation plans in response to global changes. With the rapid influx of soil biodiversity data, now is the time to take the first steps forward in establishing a global soil biodiversity information platform.

1 1. Introduction

2 Soils are increasingly recognized as crucial components of ecosystems and biodiversity
3 (Wardle et al., 2004; Bardgett and Wardle 2010), and they represent unique compartments of
4 terrestrial ecosystems by comprising components of the atmosphere, biosphere, hydrosphere,
5 and lithosphere. Soil biodiversity supports many terrestrial ecosystem functions (Wall et al.,
6 2012) and delivers important ecosystem services such as food and fiber production, carbon
7 sequestration, and degradation of pollutants (Wall et al., 2010; Wardle 2002). However, the data
8 and information regarding diversity that lives in soil remains insufficiently catalogued and
9 coordinated, and this limits our ability to fully assess the key role soil biodiversity plays in
10 supporting terrestrial systems and ecosystem services. In contrast to soil systems, greater effort
11 has been put towards cataloguing global diversity in marine and other terrestrial systems (Jetz
12 et al., 2012; Appeltans et al., 2012; Canhos et al., 2014; Hudson et al., 2014) and into making
13 these data free and open access (Guralnick et al., 2007; Wiezorek et al., 2012). Global efforts to
14 synthesize biodiversity data have proven highly successful in the transfer of information, have
15 improved our understanding of species ecology and distribution patterns, and allows for better
16 monitoring and response plans to global change effects (Dirzo et al., 2014; Hampton et al.,
17 2013). Given that we are facing unprecedented environmental alterations through climate
18 change, land use change, soil erosion, invasive species, desertification and pollution, a better
19 understanding of the global distribution and drivers of soil biodiversity is urgently needed to
20 forecast functional changes of terrestrial ecosystems and to develop appropriate management
21 practices. Therefore, here we review the rationale behind and the benefits of bringing together
22 soil biodiversity data and information through a single global data platform.

23
24 Although it is known that soils are extraordinarily diverse, the scale of soil biodiversity is not yet
25 fully understood (Wall et al., 2010). Global patterns of soil biodiversity are at most weakly
26 documented (Decaëns 2010; Tedersoo et al., 2014), and the locations of many soil biodiversity
27 hotspots have not been identified. Part and parcel to the plethora of hyperdiverse taxonomic
28 groups, global patterns of soil biodiversity are thought to differ significantly from what is reported
29 aboveground (Maraun et al., 2007; Decaëns, 2010; Ramirez et al., 2014; Tedersoo et al., 2012).
30 For example, soil microorganisms do not respond to large-scale environmental gradients in the
31 same way as metazoans and belowground biodiversity hotspots do not necessarily mirror
32 aboveground biodiversity patterns (Fierer and Jackson 2006; Wu et al., 2011) Further, many
33 species residing in soil remain taxonomically, phylogenetically, and functionally undescribed.
34 This is most notable for microorganisms (McDonald et al., 2012) but it is also true for soil fauna
35 (Bik et al., 2012; Behan-Pelletier 1999; Rougerie 2009). Therefore, categorizing species into
36 discrete taxonomic units represents a challenge for soil biodiversity documentation where many
37 of the species' characteristics and phylogenies are not yet available (Bardgett and van der
38 Putten 2014).

39
40 Regardless of these challenges, soil biodiversity research has dramatically increased over the
41 last three decades, and the scope of soil biodiversity data is immense. Soil biodiversity data
42 types range from classical specimen based collections (Burkhardt et al., 2014) to molecular and
43 genomics samples (Gilbert et al., 2014). In between are a wide spectrum of community-
44 aggregated data (i.e. trophic levels to relative abundances) organism attributes (e.g.
45 abundance, biomass and traits), and environmental measurements (e.g. georeference
46 coordinates, biome type, soil characteristics and climatic variables). Like other biodiversity
47 information, soil biodiversity data can be digital and available online, though much data remains
48 'dark' – not digitized or not available (Heidorn 2008). Whether in a national repository, stored on
49 a personal computer, or found in a museum drawer, the first step in any data synthesis project
50 is to make dark data digitally accessible [Box 1] (Hill et al., 2012). Next is to establish a
51 mechanism to link digitally available data globally (such as an online portal).

53 Here we present an independent initiative to assess and store information on global soil
54 biodiversity; to link species, environmental and other data and make data accessible at a global
55 level. Our goal was to propose a system that could be linked to other biodiversity and
56 ecosystem relevant databases, accommodate new and future methods and technologies, be
57 useful to a wide array of end users (from the public to scientists to policy makers), and be free
58 and open access.
59

60 **BOX 1:** *Digital soil biodiversity information is currently stored in a wide array of databases,*
61 *warehouses, catalogues and other repositories, and contains various types of data (see*
62 *Supp. Table 1 for a more extensive list of examples).*

- 63 • **Catalogues:** *Taxonomy lists with descriptions of the organism. May have occurrence*
64 *data and may contain images, videos or other media. (Example: Encyclopedia of*
65 *Life)*
- 66 • **Data Warehouse:** *An information system that links taxonomy (morphology and/or*
67 *annotated sequences) and ecological information across databases and individual*
68 *studies. (Example: Edaphobase)*
- 69 • **Public or Private Databases:** *Species lists for a given study, experiment or location.*
70 *May include any number of additional measured parameters such as soil*
71 *environment measurements and climate information. (Example: Earth Microbiome*
72 *Project)*
- 73 • **Sequence Archives:** *Nucleotide sequences that provide valuable information on*
74 *relevant organisms. These can be useful for determining phylogenies and functional*
75 *characteristics of organisms. May follow standards of Genomic Standards Consortia.*
76 *(Example: European Nucleotide Archive (ENA), National Center for Biotechnology*
77 *Information (NCBI))*

78 **Applied Advances**

79 It is now commonplace to concurrently survey soil biodiversity and explore the role these
80 organisms play in ecosystem functions and global sustainability (Wall et al., 2012; Bardgett and
81 van der Putten 2014). However, we still lack baseline values for soil biodiversity as well as
82 reference values (either abundance ranges or occurrence) that may prove critical in assessing
83 the current status of soils and implementing management and policy efforts to keeping soils and
84 soil biodiversity in a so-called 'normal operating range' (Koch et al., 2013; Jackson et al., 2007).
85 This will be particularly important as we continue to understand the impact of certain global
86 changes on soil biodiversity and their interactions within functioning food webs (Blankinship et
87 al., 2011; Garcia et al., 2014). For example, agricultural intensification reduces the abundance
88 of soil fungi relative to bacteria, reduces earthworms, mycorrhizal fungi, and increases the
89 numbers of plant parasitic nematodes (Tsiafouli et al., 2015). Less is known on effects of
90 incipient changes, or changes that encompass temporally complex and indirect feedback
91 effects, such as consequences of global warming, biological invasions, or habitat fragmentation
92 (Dickie et al., 2014; Blankinship et al., 2011; Lindo et al., 2012).
93

94 Reference values can be an important tool for determining the success of ecosystem restoration
95 and comparing data across time scales (Frouz; et al., 2004; Kardol and Wardle 2012) and for
96 detecting subtle trends in temporal soil biodiversity assessments (Bardgett 2005). Specific
97 indicators that can be accessed from a global platform, such as disease-suppression (Mendes
98 et al., 2011) and nutrient retention capacity of soil (de Vries et al., 2013), can also be used by
99 land managers in order to calibrate and further improve sustainability of production methods, or
100 used to develop rapid and economic soil biodiversity assessment tools for use by policy makers
101 and end users (Bone et al., 2014; Wall et al., 2012). As demonstrated by the Global Biodiversity
102

103 Information Facility and other global data synthesis efforts (Otegui et al., 2013), access and
104 availability of data has helped to predict the impact of climate change (Warren et al., 2013),
105 monitor invasive species (Gatto et al., 2013) and inform on issues like human health (Daszak et
106 al., 2013) and food and farming (Vincent et al., 2013). Further, the efforts by GBIF and Map of
107 Life (MOL) support the work of the CBD, IPBES, GEO-BON, and many others (see GBIF.org).
108 The inclusion of soil biodiversity data in such global assessments is a highly important and
109 necessary next step.

110

111 **Theoretical and research advances**

112 The prospect of accessing global soil biodiversity information through a single portal will create
113 novel opportunities to develop, refine and test underlying ecological theory. The synthesis of
114 biodiversity data across larger spatial scales and greater taxonomic breadth may uncover
115 emergent properties that cannot currently be foreseen (Brose et al., 2012) and will give better
116 insight into species' ecological preferences and geographical ranges (Brose et al., 2004;
117 Tedersoo 2014; Fierer et al. 2013). Here we identify five topic areas that, while not exhaustive,
118 will be enhanced by a global data platform effort:

119

120 (1) *Macroecology and biogeographical patterns*: Characterizing global patterns is of paramount
121 importance for conservation of soil biodiversity and global change scenarios on the
122 functioning of soil systems in a future world. A comprehensive view of biogeographic
123 patterns will be critical to reveal important scientific questions, to discover where and why
124 there are hot spots of biodiversity, to identify the drivers of belowground diversity, and will
125 ultimately boost the use of macroecological approaches in soil ecology research (Fierer et
126 al., 2013; Tedersoo et al., 2014).

127 (2) *Biodiversity maintenance and loss*: A synthesis of soil biodiversity data will help identify
128 drivers and mechanisms underlying both the maintenance and loss of biodiversity in soil and
129 dependent terrestrial systems. The support that belowground diversity gives to aboveground
130 diversity is drastically underestimated, and by overlaying belowground and aboveground
131 biodiversity patterns we can better assess the impact of biodiversity losses. Further, these
132 efforts may prove especially important in terms of invasion ecology, identifying which groups
133 are prone to invade (e.g. earthworms (Hendrix et al., 2008)), and the mechanisms facilitating
134 invasion (e.g. Dickie et al., 2014) and prevention efforts.

135 (3) *Ecosystem functions and services*: Soil organisms co-determine a plethora of provisioning
136 and regulating ecosystem services (Wardle et al., 2004; Lavelle et al., 2006), but the
137 appreciation of their functional significance remains deficient due to their cryptic nature and
138 overlapping functions (Setälä et al., 2005). While conventional anthropogenic land
139 management practices often have aimed to optimize certain (single) ecosystem functions or
140 services (Cardinale et al., 2012), soil biodiversity exemplifies the value of multifunctional
141 ecosystems (Wagg et al., 2014; Setälä et al., 2014). Recent evidence shows that the
142 structure and composition of the soil community and the presence of specific functional
143 groups, is key to delivering a range of ecosystem services, such as N retention and C
144 storage (de Vries et al., 2013; Lange et al., 2015).

145 (4) *Community ecology*: Soil communities are notoriously complex and conventional community
146 ecological theory may be challenged by the spatially complex habitat soil organisms live in
147 (Ettema and Wardle 2002). Multitrophic soil biodiversity assessment may help to refine
148 existing soil food web models (Digel et al., 2014). Further, global-scale information on the
149 co-occurrence of different taxa in soil will shed light on the relative significance of trophic vs.
150 non-trophic interactions in soil, top-down vs bottom up forces and their interplays (Moore et
151 al., 2004) and ecological network perspectives may provide useful tools to clarify
152 interactions among the different soil functional groups and to certain ecosystem functions
153 (Barberán et al., 2011; Morriën and van der Putten 2013).

154 (5) *Aboveground-belowground interactions*: As our knowledge of belowground communities
155 increases, so too does our awareness of the important, complex interactions between soil
156 organisms and aboveground biodiversity (Hooper et al., 2000). By revealing belowground
157 biodiversity patterns, we can gain better insight into the linkages between above- and
158 belowground systems. Plus, soil biodiversity data will be made more valuable if it can be
159 clearly linked to with data pertaining to aboveground communities (such as through the Map
160 of Life or GBIF).
161

162 **A proposed framework**

163 Our ability to address a range of applied and theoretical questions, or to assess biogeographical
164 patterns, is to a large extent limited by access and integration of the available data. Currently,
165 there is no single repository or platform that allows access to soil biodiversity information,
166 across all species, or at a global scale. Therefore, we propose a framework to initiate linking
167 different databases and repositories via the internet (Fig 1). The end platform will be both a
168 database and a free, open access portal to link various national and local data sources around
169 the world. Linking data from existing databases is not trivial, nor is it a new challenge (Jetz et
170 al., 2012). Previous efforts such as GBIF and MOL have demonstrated that because there are
171 no required guidelines or consistency between studies or pre-established databases, minimum
172 standards and classifications must be identified. Soil biodiversity standards must then be
173 harmonized with the global standards already in place (e.g. Wiczorek et al., 2012; Yilmaz et al.,
174 2011). While applying even simple standards will lead to the omission of some studies and data,
175 quality of the data will be valued over quantity, ultimately resulting in a higher quality synthesis.
176

177 Integration and access to soil biodiversity data will be accomplished in three phases: discovery,
178 standardization and a final user interface:
179

180 **Phase I - “Discover” where soil biodiversity data is housed**: This phase will be two-fold;
181 first to establish a taxonomy list - a list of organisms living in the soil, and second to
182 inventory soil biodiversity information. The taxonomy list will be shared with the Global
183 Biodiversity Information Facility (GBIF) to tag preexisting soil related biological
184 observations that can thereon be searched and queried (much like the Global Mountain
185 Biodiversity Assessment (GMBA) (gmba.unibas.ch)) and allow for easier integration of
186 new data. The ‘taxonomy list’ and an inventory of soil biodiversity information will be
187 made available through the Global Soil Biodiversity Initiative (GSBI). It is in this stage
188 that data quality will be also assessed, a complicated issue all biodiversity data studies
189 must deal with. We propose to follow guidelines set forth and established by GBIF.
190

191 **Phase II - Establish a standardization framework by which to link past, present and
192 future data**: Besides taxonomic synonyms it also will be necessary to develop and
193 implement thesauri for the various information fields (i.e. regarding habitat or climate
194 parameters, methods etc.). Standardized ontologies are necessary to link between
195 different data sources and into GBIF (Supp. 1) and other global data centers (such as
196 MOL, ISRIC, EOL, Genbank and others). Furthermore, to allow data comparability from
197 the individual data sources, standardization of numeric (abundances, pH values, etc.)
198 and nominal (i.e., habitat types, soil types) data will be crucial. Concurrently, we must
199 also establish the minimum set of parameters needed, and formalize data copyright
200 privacy and licensing rules. Together these efforts will provide the critical foundation and
201 quality criteria on which to build the platform.
202

203 *Short read sequence data: In the case of microbial marker gene sequence data
204 (either 16S, 18S, ITS or similar) it is difficult to extract taxonomic information for a*

205 *number of reasons (otu picking methods, chimeras, read length, Orgiazzi et al.,*
206 *2014). Plus due to the enormous amount of sequence data, reprocessing the full*
207 *datasets would not be tractable. Therefore, we propose to link short read*
208 *sequence data by location, rather than by taxon identification. This is based on*
209 *the fact that there is currently no consensus on the correct protocol for handling*
210 *these data, and integrating processed sequence data would introduce substantial*
211 *methodological artifacts (Caporaso et al., 2010). Instead, our approach allows*
212 *convenient access to these data linked to geography and allows users to process*
213 *the data of interest using a consistent protocol based on individual research*
214 *questions.*

215
216 **Phase III - Establish a user-friendly interface that allows for the integration and**
217 **comparison of soil biodiversity data - here called 'Soil Portal':** The portal will be designed
218 specifically for manipulation and analyses of the data in order to address the theoretical
219 questions outlined above and to provide stakeholders with the type of information
220 needed for management and policy decisions. It is in this phase that we would finally be
221 able to combine collection data across taxonomic groups, spatial scales and research
222 experiments. As demonstrated previously (Hill et al. 2012), users are reluctant to use
223 any interface that costs time, therefore, we propose a platform that would offer
224 researchers a set of tools, rewards for contributing their data to the community- such as
225 data analyses tools, DOIs for data publication, and a link to other initiatives and data
226 portals.

227 228 229 **Outlook**

230 In order to progress this project, first, buy-in from the community of soil biologists is required;
231 our goal is to galvanize and guide soil ecologists to make their data available. Researchers can
232 continue to upload data from their home repositories, data will not have to be uploaded more
233 than once, and there is no need to support a single, comprehensive database - a monetarily
234 expensive and time consuming task. The framework is designed so that participation in the
235 effort to liberate individual datasets will only require minor changes to how researchers work
236 (i.e. time for data input and training for students and young scientists), but has the potential for
237 great individual rewards such as more publications (e.g. 'data papers'), increased exposure
238 leading to invitations and collaborations, as well as reciprocal access to a wealth of data from
239 colleagues. Admittedly, in addition to the technical challenges outlined in the introduction, the
240 main limiting factor of this proposal will be resources. Specifically, time and funds must be
241 invested upfront to move this effort forward in an efficient way.

242 243 **Conclusion**

244 In response to unprecedented global environmental changes and the drastic impacts on
245 biodiversity (Sala et al., 2000), there is a sense of urgency to bring together global biodiversity
246 information that will provide the basis to determine the species and communities that are
247 particularly vulnerable to change and extinctions (Jetz et al. 2012; Cardinale et al 2012; Scholes
248 2008) and focus conservation and management practices (Turner et al., 2015). The organisms
249 that live in the soil are no exceptions. The focus of the outlined framework goes beyond species
250 information, and therefore a major challenge and goal will be to integrate the different
251 information types whereby a range of ecological questions can be addressed. Soil biodiversity
252 information is of broad interest to other disciplines, including plant ecologists, agriculturalists,
253 invertebrate ecologists, carbon and climate modelers, and would open new unique opportunities
254 for collaboration between the groups. As such, we have designed a framework that will interface
255 with other disciplines through GBIF and the like. In addition to data access and standardization,

256 a priority of this effort will be analytical and visualization tools for end users. Beyond progressing
257 scientific research these tools should help to communicate results and bring the interest of a
258 larger, more general audience. All together, access to rapidly accumulating soil biodiversity
259 information across the globe has the potential to improve research and elevate soil ecology to
260 be on par with our understanding of aboveground systems.

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502 **Figures:**

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504 **Figure 1:** Integration and access to soil biodiversity data will be accomplished in three phases:
505 (I) discovery, (II) standardization and (III) a final user interface, and the timing of these phases
506 will be directly related to the effort and resources put in to the framework.

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Figure 1.JPEG

