

Advancing subterranean conservation through Global Research on eDNA in Groundwaters (GReG)

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If water is life, then groundwater is its lifeblood—essential, but hidden, it forms the largest reservoir of unfrozen freshwater globally. Populated by highly specialized, often relictual lineages across the tree of life, groundwater ecosystems not only support remarkable taxonomic, functional and phylogenetic diversity (Griebler and Lueders 2009; Martínez et al. 2018; Marmonier et al. 2023), but also play a crucial role in sustaining the functioning of the global water cycle (Griebler and Avramov 2015; Saccò et al. 2024). Consequently, the protection of groundwaters as ecosystems, which can be affected by overextraction, warming or pollution, is essential. Despite this, groundwater is largely absent from policy frameworks and overlooked by large initiatives such as the European Water Framework Directive (Di Lorenzo et al. 2024) that are designed to protect the environment (Fiser et al. 2022, 2025; Griebler et al. 2023). While the human impact on above-ground global biodiversity has triggered widespread conservation responses, subterranean ecosystems continue to suffer from a lack of visibility, research attention, and regulation (Saccò et al., 2025a). Current global knowledge on groundwater biodiversity and ecological dynamics is far from complete, hindering effective assessment and conservation efforts (Mammola et al. 2024). Regrettably, groundwater species are facing escalating cumulative threats including climate and land use change, contamination, over-extraction, heating and salinization, amongst many others (Manenti et al. 2021; Becher et al. 2022; Couton et al. 2023a; Nanni et al. 2023; Vaccarelli et al. 2023). Left unaddressed, these threats compromise not only biodiversity but also the vast number of services that groundwater ecosystems unnoticeably provide to both nature and human society (Boulton et al. 2008; Griebler and Avramov 2015; Mammola et al. 2025; Saccò et al. 2025b).

The development of conservation strategies for groundwater and groundwater-dependent ecosystems (GDE, ecosystems that rely on groundwater for some or all of their water needs) requires an improved understanding of subterranean ecological dynamics, particularly under ongoing climatic changes (Boulton et al. 2023; Griebler et al. 2023; Vaccarelli et al. 2023). To address these challenges, increased collaborative efforts based on new technologies, scientific rigour, interdisciplinarity and inclusivity are required (Saccò et al. 2025b). In this regard, the analysis of environmental DNA (eDNA) is a technique proving to be an effective bridge connecting researchers,

stakeholders and citizens over a myriad of aquatic ecosystems (Deiner et al. 2017; Takahashi et al. 2023; Blackman et al. 2024). Broadly defined as the total pool of DNA isolated from environmental samples (Pawlowski et al. 2020), eDNA has been successful at unveiling biodiversity patterns from regional (Nakagawa et al. 2018) to global scales (Ficetola et al. 2024), and has been applied in taxonomy-free (Apothélos-Perret-Gentil et al. 2017) and functional (Condachou et al. 2023; Cantera et al. 2025) approaches. Moreover, eDNA is being increasingly combined with other large-scale monitoring tools (e.g., satellite data, Zong et al. 2024) and broader biodiversity datasets (Abarenkov et al. 2023; Takahashi et al. 2025) to advance a rich, diverse research agenda. Concerning groundwater, eDNA research is gaining similar momentum (e.g., Koch et al. 2022; Saccò et al. 2022; Couton et al. 2023a, b; Al-Rashidi et al. 2025; Guzik et al. 2025a, b), despite a lag in protocol validation, application, and DNA-assay development compared to other biomes (e.g., oceans or rivers - Altermatt et al. 2025).

With the intention of disrupting the current inertia in the global conservation of groundwater ecosystems - where many calls for action have gone unheeded (e.g., Sánchez-Fernández et al. 2021; Wynnee et al. 2021; Di Lorenzo et al. 2024) - we present here a research initiative defined by the acronym of GReG: Global Research on eDNA in Groundwaters. The first pitch for this collaborative global eDNA study in groundwater ecosystems was proposed during the 26th International Conference on Subterranean Biology (ICSB) held in Sardinia in September 2024. Nine months later, on the 5th of June 2025, GReG was launched *via* an online seminar, led by Dr Mattia Saccò, Dr Michelle Guzik and Dr Kathryn Korbel. A gathering of 56 representative researchers and regulators across the globe took place, and guest speakers provided insights regarding global eDNA studies in oceans (Citizens of the Sea; Pochon et al. 2025), open and FAIR (Findable, Accessible, Interoperable and Reusable) data practices for sharing eDNA data (Takahashi et al. 2025), taxon-free eDNA approaches in groundwater studies, and how to utilize eDNA to address global biodiversity targets (Altermatt et al. 2025). The second part of the seminar was dedicated to the planning of a self-funded project in collaboration with partners such as Biodiversa+ projects (DarCO - Mammola et al. 2024; Sub-BioMon - Zgamałster et al. 2024), research laboratories (TrEnD Lab - <https://research.curtin.edu.au/scieng/research/trend-lab/>; Altermatt Lab - <https://www.altermattlab.ch/>) and global datasets (Stygofauna Mundi - Martínez et al. 2018).

Planned to be carried out in two phases, GReG is conceived as a unifying platform for researchers and diverse stakeholders from government and industry to meet and significantly advance groundwater science primarily through eDNA and genomics. Phase 1 (June 2025–June 2026) will involve fieldwork at over 50 well-known sites (karst, alluviums and anchialine caves) across four continents, using standardized sampling and analytical protocols targeting biodiversity across the entire Tree of Life. In addition, a survey on methodological approaches in groundwater eDNA research has been circulated to all the collaborators, aimed at understanding current practices and limitations for eDNA analysis in groundwater environments. Further projected research outputs include peer-reviewed papers on groundwater

eDNA protocols and publicly available DNA barcode quality assessments for stygofauna. Phase 2 (from July 2026) will build on the lessons learned from Phase 1 to expand the project to the Global South and other under-investigated areas of the world *via* participatory science approaches. This Forum Paper also serves as an opportunity to make a public call to the researchers and stakeholders from those regions, with an ambition of widening novel groundwater-based research beyond well-known global biodiversity hotspots. The outcomes derived from this collaborative effort will involve scientific publications, data sharing following the FAIR eDNA guidelines, workshops and media articles, with the objective to advance subterranean aquatic eDNA research. Last but not least, GReG also aims to raise awareness regarding the importance of groundwater ecosystems - a notoriously arduous task for an out of sight, out of mind environment (Boulton et al. 2023).

A shared vision amongst GReG collaborators is that the key to effectively preserving unique subterranean biota while using groundwater resources sustainably is not yet lost; we just need the right team of locksmiths (researchers, regulators, stakeholders, citizens) to come together and cut the right set of biomonitoring keys and conservation guidelines. GReG and its multidisciplinary approach provide the first step towards preserving subterranean biota at the global level, and thus utilising groundwater resources sustainably—an ambitious but vital goal for future water security.

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References

- Altermatt F, Couton M, Carraro L, Keck F, Lawson-Handley L, Leese F, Xiaowei Z, Zhang Y, Blackman RC (2025) Utilizing aquatic environmental DNA to address global biodiversity targets. *Nature Reviews Biodiversity* 1: 332–346. <https://doi.org/10.1038/s44358-025-00044-x>
- Al-Rashidi A, Sabarathinam C, Samayamanthula DR, Alsabti B, Habibi N, Manickam N (2025) Environmental DNA signatures as a tool to trace the groundwater contamination mechanisms and its associated biodiversity; Applications, limitations and future directions. *Current Opinion in Environmental Science & Health* 45: 100622. <https://doi.org/10.1016/j.coesh.2025.100622>
- Andersson AF, Bissett A, Finstad AG, Fossøy F, Grosjean M, Hope M, Jeppesen TS, Kóljalg U, Lundin D, Nilsson RH, Prager M, Svenningsen C, Schigel D (2021) Publishing DNA-derived data through biodiversity data platforms. Version 71ae0fa.
- Apothéloz-Perret-Gentil L, Cordonier A, Straub F, Iseli J, Esling P, Pawlowski J (2017) Taxonomy-free molecular diatom index for high-throughput eDNA biomonitoring. *Molecular Ecology Resources* 17(6): 1231–1242. <https://doi.org/10.1111/1755-0998.12668>
- Becher J, Englisch C, Griebler C, Bayer P (2022) Groundwater fauna downtown - Drivers, impacts and implications for subsurface ecosystems in urban areas. *Journal of Contaminant Hydrology* 248: 104021. <https://doi.org/10.1016/j.jconhyd.2022.104021>
- Blackman R, Couton M, Keck F, Kirschner D, Carraro L, Cereghetti E, Perrelet K, Bossart R, Brantschen J, Zhang Y, Altermatt F (2024) Environmental DNA: The next chapter. *Molecular Ecology* 33(11): e17355. <https://doi.org/10.1111/mec.17355>
- Boulton AJ, Fenwick GD, Hancock PJ, Harvey MS (2008) Biodiversity, functional roles and ecosystem services of groundwater invertebrates. *Invertebrate Systematics* 22: 103–116. <https://doi.org/10.1071/IS07024>
- Boulton AJ, Bichuette ME, Korbel K, Stoch F, Niemiller ML, Hose, GC, Linke S (2023) Recent concepts and approaches for conserving groundwater biodiversity. *Groundwater ecology and evolution* (Netherlands: Academic Press), 525–550. <https://doi.org/10.1016/B978-0-12-819119-4.00001-9>
- Cantera I, Giachello S, Münkemüller T, Caccianiga M, Gobbi M, Losapio G, Marta S, Valle B, Zawierucha K, Thuiller W, Ficetola GF (2025) Describing functional diversity of communities from environmental DNA. *Trends in Ecology & Evolution* 40(2): 170–179. <https://doi.org/10.1016/j.tree.2024.10.007>
- Condachou C, Milhau T, Murienne J, Brosse S, Villéger S, Valentini A, Dejean T, Mouillot D (2023) Inferring functional diversity from environmental DNA metabarcoding. *Environmental DNA* 5(5): 934–944. <https://doi.org/10.1002/edn3.391>
- Couton M, Hürlemann S, Studer A, Alther R, Altermatt F (2023a) Groundwater environmental DNA metabarcoding reveals hidden diversity and reflects land-use and geology. *Molecular Ecology* 32(13): 3497–3512. <https://doi.org/10.1111/mec.16955>
- Couton M, Studer A, Hürlemann S, Locher N, Knüsel M, Alther R, Altermatt F (2023b) Integrating citizen science and environmental DNA metabarcoding to study biodiversity of groundwater amphipods in Switzerland. *Scientific Reports* 13(1): 18097. <https://doi.org/10.1038/s41598-023-44908-8>

- Deiner K, Bik HM, Mächler E, Seymour M, Lacoursière-Roussel A, Altermatt F, Creer S, Bista I, Lodge DM, de Vere N, Pfrender ME, Bernatchez L (2017) Environmental DNA metabarcoding: Transforming how we survey animal and plant communities. *Molecular ecology* 26(21): 5872–5895. <https://doi.org/10.1111/mec.14350>
- Di Lorenzo T, Lunghi E, Aanei C, Altermatt F, Alther R, Amorim IR, Băncilă RI, Bellvert, A, Blomberg A, Borges PAV, Brad T, Brancelj A, Brankovits D, Cardoso P, Cerasoli F, Chauveau CA, Crespo L, Csader M, Delić T, Di Cicco M, Douady CJ, Duchemin L, Faille A, Fiasca B, Fišer C, Flot J-F, Gabriel R, Galassi DMP, Garzoli L, Griebler C, Karwautz C, Keneski MI, Konecny-Dupré L, Lilley T, Malard F, Martínez A, Meierhofer MB, Messana G, Millán A, Mizerakis V, Mori N, Nanni V, Nicolosi G, Oromí P, Pallarés S, Pereira F, Reboleira ASPs, Saccò M, Salussolia A, Sánchez-Fernández D, Sarbu SM, Stefan A, Stoch F, Tabilio di Camillo A, Taiti S, Vaccarelli I, Valanne V, Zgamaister M, Zakšek V, Zittra C, Mammola, S (2024) EU needs groundwater ecosystems guidelines. *Science* 386(6726): 1103–1103. <https://doi.org/10.1126/science.ads8140>
- Ficetola GF, Marta S, Guerrieri A, Cantera I, Bonin A, Cauvy-Fraunié S, Ambrosini R, Caccianiga M, Anthelme F, Azzoni RS, Almond P, Alviz Gazitúa P, Ceballos Lievano JL, Chand P, Chand Sharma M, Clague JJ, Alejo Cochachín Rapre J, Compostella C, Cruz Encarnación R, Dangles O, Deline P, eger A, Erokhin S, Franzetti A, Gielly L, Gili F, Gobbi M, Hågvær S, Kaufmann R, Khedim N, Meneses RI, Morales-Martínez MA, Peyre G, Pittino F, Proietto A, Rabatel A, Sieron K, Tielidze L, Urseitova N, Yang Y, Zaginaev V, Zerboni A, Zimmer A, Diolaiuti GA, TAberlet P, Poulenard J, Fontaneto D, Thuiller W, Carteron, A (2024) The development of terrestrial ecosystems emerging after glacier retreat. *Nature* 632(8024): 336–342. <https://doi.org/10.1038/s41586-024-07778-2>
- Fišer C, Zgamaister M, Jemec Kokalj A, Mali N, Šumrada T, Glavan M, Hose GC, Schwartz B, Di Lorenzo T, Griebler C, Cvejić R (2025) Toward sustainable irrigation practices safeguarding groundwater biodiversity and ecosystem services. *Bioscience* 2025: biaf016. <https://doi.org/10.1093/biosci/biaf016>
- Griebler C, Lueders T (2009) Microbial biodiversity in groundwater ecosystems. *Freshwater biology* 54(4): 649–677. <https://doi.org/10.1111/j.1365-2427.2008.02013.x>
- Griebler C, Avramov M (2015) Groundwater ecosystem services: a review. *Freshwater Science* 34(1): 355–367. <https://doi.org/10.1086/679903>
- Griebler C, Hahn HJ, Mammola S, Niemiller ML, Weaver L, Saccò M, Bichuette ME, Hose GC (2023) Legal frameworks for the conservation and sustainable management of groundwater ecosystems. *Groundwater ecology and evolution* (Netherlands: Academic Press), 551–571. <https://doi.org/10.1016/B978-0-12-819119-4.15005-X>
- Guzik MT, Stringer DN, Thornhill J, Coates PJ, van der Heyde M, Hillyer MJ, white NE, Saccò M, Beasley-Hall P, Humphreys WF, Harvey MS, Huey JA, Wilson NG, Alexander J, Humphreys G, King RA, Cooper SJB, Pinder A, Perina G, Hosie AM, Kirkendale L, Nevill P, Austin AD (2025a) What are the best practices for curating eDNA custom barcode reference libraries? A case study using Australian subterranean fauna. *Biological Journal of the Linnean Society*. [in press]

- Guzik MT, Thornhill J, van der Heyde M, Stringer DN, White NE, Saccò M, Beasley-Hall P, Nevill P, King RA, Cooper SJB, Austin AD (2025b) Towards a global barcode reference library for subterranean fauna. *Science of the Total Environment*. [in press]
- Koch F, Blum P, Korbel K, Menberg K (2024) Global overview on groundwater fauna. *Ecohydrology* 17(1): e2607. <https://doi.org/10.1002/eco.2607>
- Mammola S, Altermatt F, Alther R, Amorim IR, Băncilă RI, Borges PA, Brad T, Brankovits D, Cardoso, P, Cerasoli F, Chauveau CA, Delić T, Di Lorenzo T, Faille A, Fišer C, Flot J-F, Gabriel R, Galassi DMP, Garzoli L, Griebler C, Konecny-Dupré L, Martínez A, Mori N, Nanni V, Ogorelec Z, Pallarés S, Salussolia A, Saccò M, Stoch F, Vaccarelli I, Zagmajster M, Zittra C, Meierhofer MB, Sánchez-Fernández D, Malard F (2024) Perspectives and pitfalls in preserving subterranean biodiversity through protected areas. *npj Biodiversity* 3(1): 2. <https://doi.org/10.1038/s44185-023-00035-1>
- Mammola S, Brankovits D, Di Lorenzo T, Amorim I R, Bancela R I, Bellvert A, Bernard E, Blomberg AS, Borges PAV, Cappelletti M, Ferreira R, Gabriel R, Galassi DMP, Garzoli L, Gerovasileiou V, Hose GC, Korbel KL, Mabrouki Y, Martino S, Miller AZ, Mori N, Nanni V, Nicolosi G, Saccò M, Sakihara TS, Sousa Silva M, Tamalavage A, Zagmajster M, Chávez E, Griebler C, Cardoso P, Martínez A (2025) Subterranean environments contribute to three-quarters of classified ecosystem services. *EcoEvoRxiv*. <https://doi.org/10.32942/X2JM0S>
- Manenti R, Piazza B, Zhao Y, Padoa Schioppa E, Lunghi E (2021) Conservation studies on groundwaters' pollution: challenges and perspectives for stygofauna communities. *Sustainability* 13(13): 7030. <https://doi.org/10.3390/su13137030>
- Marmonier P, Galassi DMP, Korbel K, Close M, Detry T, Karwautz C (2023) Groundwater biodiversity and constraints to biological distribution. *Groundwater ecology and evolution* (Netherlands: Academic Press), 113–140.
- Martinez A, Anicic N, Calvaruso S, Sanchez N, Puppieni L, Sforzi T, Zaupa S, Alvarez F, Brankovits D, Gašiorowski L, Gerovasileiou V, Gonzalez BC, Humphreys WH, Iliffe TM, Worsaae K, Bailly N, Fontaneto D (2018) A new insight into the Stygofauna Mundi: Assembling a global dataset for aquatic fauna in subterranean environments. *ARPHA Conference Abstracts 1*: e29514. [Pensoft Publishers] <https://doi.org/10.1016/B978-0-12-819119-4.00003-2>
- Nakagawa H, Yamamoto S, Sato Y, Sado T, Minamoto T, Miya M (2018) Comparing local- and regional-scale estimations of the diversity of stream fish using eDNA metabarcoding and conventional observation methods. *Freshwater Biology* 63(6): 569–580. <https://doi.org/10.1111/fwb.13094>
- Nanni V, Piano E, Cardoso P, Isaia M, Mammola S (2023) An expert-based global assessment of threats and conservation measures for subterranean ecosystems. *Biological Conservation* 283: 110136. <https://doi.org/10.1016/j.biocon.2023.110136>
- Pawlowski J, Apothéloz-Perret-Gentil L, Altermatt F (2020) Environmental DNA: What's behind the term? Clarifying the terminology and recommendations for its future use in bio-monitoring. *Molecular Ecology* 29(22): 4258–4264. <https://doi.org/10.1111/mec.15643>
- Pochon X, Bomati E, Frankham M, Frankham J, Swale E, Laroche O, Victor B, Stone K, Saenz P (2025) Citizens Of The Sea: Mapping Ocean Health At Scale (No. OOS2025-207). *Copernicus Meetings*. <https://doi.org/10.5194/oos2025-207>

- Saccò M, Guzik M, van der Heyde M, Nevill P, Cooper SJB, Austin AD, Coates PJ, Allentoft M, White NE (2022) eDNA in subterranean ecosystems: Applications, technical aspects, and future prospects. *Science of the Total Environment* 820: 153223. <https://doi.org/10.1016/j.scitotenv.2022.153223>
- Saccò M, Huggins X, Martínez A, Reinecke R (2025) Collaborative Science for Groundwater Biodiversity Conservation. *Ground Water* 63(4): 450–451. <https://doi.org/10.1111/gwat.13495>
- Saccò M, Mammola S, Altermatt F, Alther R, Bolpagni R, Brancelj A, Brankovits D, Fišer C, Gerovasileiou V, Griebler C, Guareschi S, Hose GC, Korbel K, Lictevout E, Malard F, Martínez A, Niemiller ML, Robertson A, Tanalgo KC, Bichuette ME, Borko S, Brad T, Campbell MA, Cardoso P, Celico F, Cooper SJB, Culver D, Di Lorenzo T, Galassi DMP, Guzik MT, Hartland A, Humphreys WF, Lopes Ferreira R, Lunghi E, Nizzoli D, Perina G, Raghavan R, Richards Z, Reboleira ASPs, Rohde MM, Sánchez Fernández D, Schmidt SI, van der Heyde M, Weaver L, White NE, Zagamajster M, Hogg I, Ruhi A, Gagnon MM, Allentoft ME, Reinecke R (2024) Groundwater is a hidden global keystone ecosystem. *Global change biology* 30(1): e17066. <https://doi.org/10.1111/gcb.17066>
- Saccò M, Allan C, Griebler C, Mammola S, Moock C, Mitchell M, Reinecke R (2025) Groundwater ecosystem services: social-ecological insights. *Groundwater Pollution and Integrated Concepts for Enhanced Knowledge and Management* (Elsevier). [in press]
- Sánchez-Fernández D, Galassi DM, Wynne JJ, Cardoso P, Mammola S (2021) Don't forget subterranean ecosystems in climate change agendas. *Nature Climate Change* 11(6): 458–459. <https://doi.org/10.1038/s41558-021-01057-y>
- Takahashi M, Saccò M, Kestel JH, Nester G, Campbell MA, Van Der Heyde M, Heydenrych MJ, Juszkiwicz DJ, Nevill P, Dawkins KL, Bessey C, Fernandes K, Miller H, Power M, Mousavi-Derazmahalleh M, Newton JP, White NE, Richards ZT, Allentoft ME (2023) Aquatic environmental DNA: A review of the macro-organismal biomonitoring revolution. *Science of the Total Environment* 873: 162322. <https://doi.org/10.1016/j.scitotenv.2023.162322>
- Takahashi M, Frøslev TG, Paupério J, Thalinger B, Klymus K, Helbing CC, Villacorta-Rath C, Silliman K, Thompson LR, Jungbluth SP, Yong SY, Formel S, Jenkins G, Laporte M, Deagle B, Rajbhandari S, Stjernegaard Jeppesen T, Bissett A, Jerde C, Hahn EE, Schriml LM, Hunter C, Newman P, Woollard P, Harper LR, Dunn N, West K, Haderlé R, Wilkinson S, Acharya-Patel N, Lopez MLD, Cochrane G, Berry O (2025) A Metadata Checklist and Data Formatting Guidelines to Make eDNA FAIR (Findable, Accessible, Interoperable, and Reusable). *Environmental DNA* 7(3): e70100. <https://doi.org/10.1002/edn3.70100>
- Vaccarelli I, Colado R, Pallares S, Galassi DM, Sanchez-Fernandez D, Di Cicco M, Meierhofer MB, Piano E, Di Lorenzo T, Mammola S (2023) A global meta-analysis reveals multilevel and context-dependent effects of climate change on subterranean ecosystems. *One Earth* 6(11): 1510–1522. <https://doi.org/10.1016/j.oneear.2023.09.001>
- Zagamajster M, Balázs G, Biró A, Bucur R, Collard O, DeliĆ T, Kos A, Flot J-F, Haidau C, Herczeg G, Iepure S, Lippert S, Lunghi E, Mntoiu S, Moldovan OT, Sambor O, Sitar C, Stoch F, Zakšek V, Weber D, Weigand A, Weigand H (2024) The Sub-BioMon Project: Developing and testing approaches to monitor subterranean biodiversity in karst. *Karst Science Days Symposium Proceedings*, 51–53. <https://doi.org/10.70655/KSD.2024.15>

- Zong S, Brantschen J, Zhang X, Albouy C, Valentini A, Zhang H, Altermatt F, Pellissier L (2024) Combining environmental DNA with remote sensing variables to map fish species distributions along a large river. *Remote Sensing in Ecology and Conservation* 10(2): 220–235. <https://doi.org/10.1002/rse2.366>
- Wynne JJ, Howarth FG, Mammola S, Ferreira RL, Cardoso P, Lorenzo TD, Galassi DMP, Medellín RA, Miller BW, Sánchez-Fernández D, Bichuette ME, Biswas J, Blackeagle CW, Boonyanusith C, Maorim IR, Borges PAV, Boston PJ, Cal RN, Cheeptham N, Deharveng L, Eme D, Faille A, Fenolio D, Fišer C, Fišer Z, ‘Ohukani‘ōhi‘a Gon SM, Goudarzi F, Griebler C, Halse S, Hoch H, Kale E, Katz AD, Kováč L, Lilley TM, Manchi S, Manenti R, Martínez A, Meierhofer MB, Miller AZ, Moldovan OT, Niemiller ML, Peck SB, Pellegrini TG, Pipan T, Phillips-Lander CM, Poot C, Racey PA, Sendra A, Shear WA, Silva MS, Taiti S, Tian M, Venarsky MP, Pakarati SY, Zgmajster M, Zhao Y (2021) A conservation roadmap for the subterranean biome. *Conservation Letters* 14(5): e12834. <https://doi.org/10.1111/conl.12834>