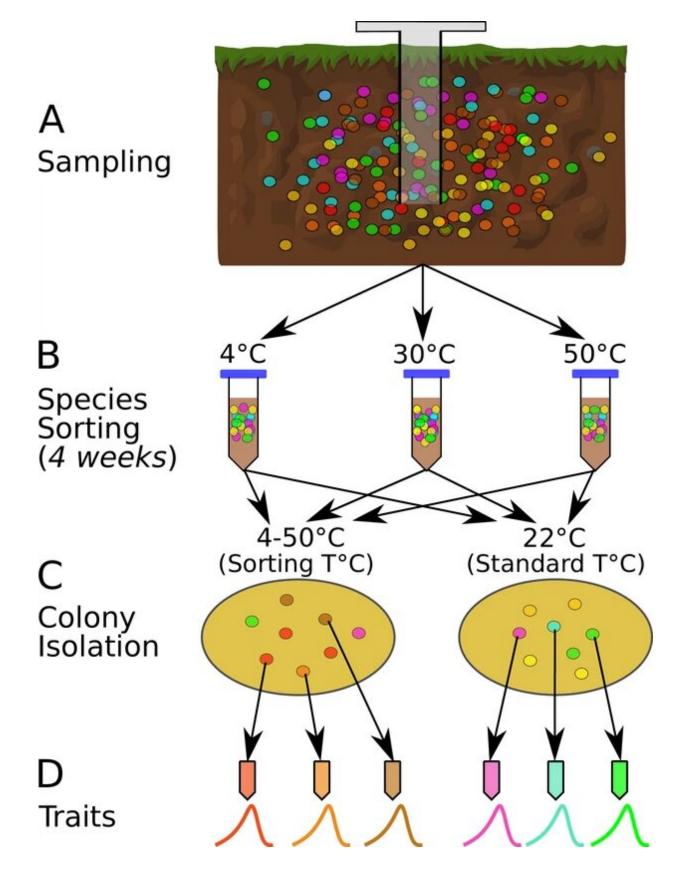


## Dormant microbes can 'switch on' to cope with climate change

November 29 2022, by Hayley Dunning







The species sorting experiment. (A) Different bacterial taxa (colored circles) sampled from the soil community. (B) Samples maintained at 4, 10, 21, 30, 40, and 50°C (only three temperatures shown for illustration), allowing species sorting for 4 weeks. (C) Soil washes from each core plated out onto agar and grown at both the sorting temperature and 22°C (standard temperature) to allow further species sorting and facilitate isolation (next step). (D) The six most abundant (morphologically different) colonies from each plate were picked, streaked, and isolated, and their physiological and life history traits measured. The curves represent each strain's unique unimodal response of growth rate to temperature. Credit: *eLife* (2022). DOI: 10.7554/eLife.80867

Dormant strains of bacteria that have previously adapted to cope with certain temperatures are switched back on during climatic change, study shows.

The results, led by a team at Imperial College London and published today in *eLife*, have important implications for predicting the impact of global warming on ecosystems.

Microbes are important for ecosystem function, because of their key roles as pathogens, <u>food sources</u> and in nutrient recycling. To understand the profound impact of climate change on the function of different ecosystems, it is therefore necessary to study the microbes within them.

Microbes live in communities of different species and strains, with each performing different functions based on their traits—characteristics such as heat tolerance and the ability to use certain chemicals.

Lead author Dr. Thomas Smith, from the Georgina Mace Center for the Living Planet at Imperial, said, "Microbial communities can respond to warming in the short term by acclimation—developing unique traits to suit the environment—or through the longer term by adaptation, where



they make evolutionary changes over many generations.

"But there is also a third mechanism, called species sorting, whereby the composition of the overall community—that is, which species are present—alters with changes in <u>temperature</u>. The importance of species sorting relative to acclimation and adaptation has not previously been explored in the context of microbial community responses to changing temperature."

## **Species sorting**

To address this, the team carried out a species sorting experiment, where they grew soil bacteria communities collected from a single site at different temperatures ranging from 4°C to 50°C.

They then measured the growth and metabolism of each isolated strain of bacteria across these different temperatures to determine their thermal performance, and studied the genetic sequences of isolated bacteria to see how temperature-response traits evolved over time.

They found that evolutionarily and functionally distinct communities emerged at each of the temperature conditions, driven by the resuscitation of microbial strains that had been inactive under previous environmental conditions.

This suggests that—rather than new bacteria moving into a community to suit the new conditions—the parent community harbors multiple bacterial strains that are pre-adapted to survive at different temperatures and can switch on when their preferred temperature is reached. As a result, microbial communities in nature are likely to be able to respond rapidly to temperature fluctuations.

Senior author Professor Thomas Bell, from the Georgina Mace Center



for the Living Planet at Imperial, said: "Understanding the relative importance of acclimation, adaptation and <u>species</u> sorting in the assembly and turnover of microbial communities is key to determining how quickly they can respond to temperature changes.

"We have found that the resuscitation of functional diversity within a microbial community can allow the whole community to survive in response to temperature changes. Further studies on other <u>microbial communities</u>—such as those residing in water—will support more accurate predictions of the effects of climate change on different ecosystems."

"Latent functional diversity may accelerate microbial community responses to temperature fluctuations," by Thomas P Smith, Shorok Mombrikotb, Emma Ransome, Dimitrios—Georgios Kontopoulos, Samraat Pawar, and Thomas Bell is published in *eLife*.

**More information:** Thomas P Smith et al, Latent functional diversity may accelerate microbial community responses to temperature fluctuations, *eLife* (2022). DOI: 10.7554/eLife.80867

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