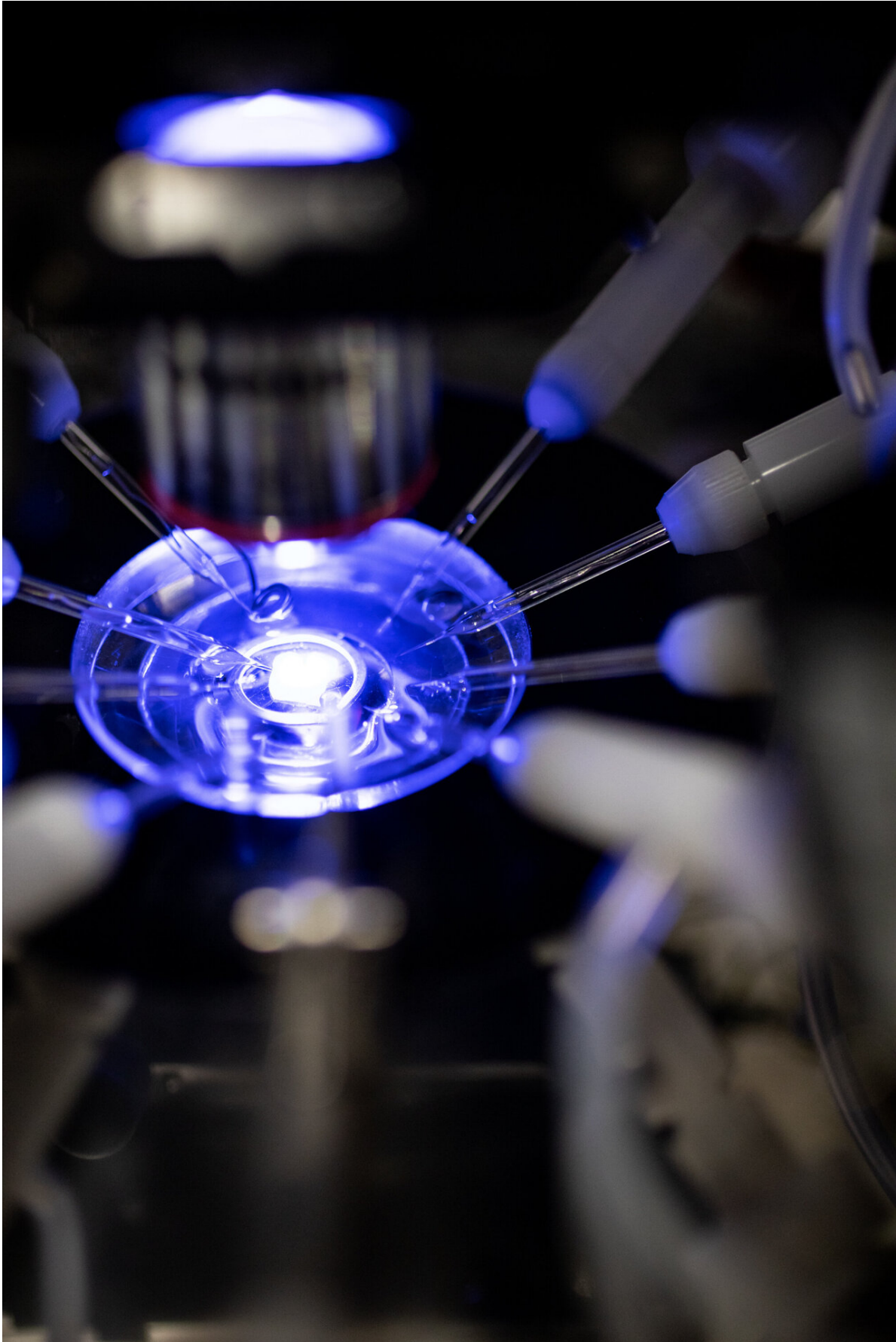


Full-scale, biologically realistic model of mouse hippocampus uncovers new mechanism for pattern separation

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Using parameters measured in the mouse hippocampus, the researchers created a comprehensive network model of the brain region. Credit: © Peter Rigaud/IST Austria

Our brains can distinguish highly similar patterns, thanks to a process called pattern separation. How our brains separate patterns is, however, not yet fully understood. Using a full-scale computer model of the dentate gyrus, a brain region involved in pattern separation, Peter Jonas, professor at the Institute of Science and Technology (IST) Austria, found that inhibitory neurons activated by one pattern suppress all their neighboring neurons, thereby switching off "competing" similar patterns. This is the result of a study published in *Nature Computational Science*.

A black cat is deceptively similar to a panther, apart from its size. Making this distinction is crucial when being pursued, and something humans achieve thanks to pattern separation, the process in which the brain distinguishes highly similar patterns and triggers very different behaviors as a response: stroke the cat or flee the panther. This process is also linked to learning. "We need to not just separate similar patterns, but also store and accurately retrieve them, for example for when we next meet a panther. Therefore, we investigated how pattern separation occurs in the hippocampus, a key memory circuit," says Peter Jonas, Professor at IST Austria and lead author of the study.

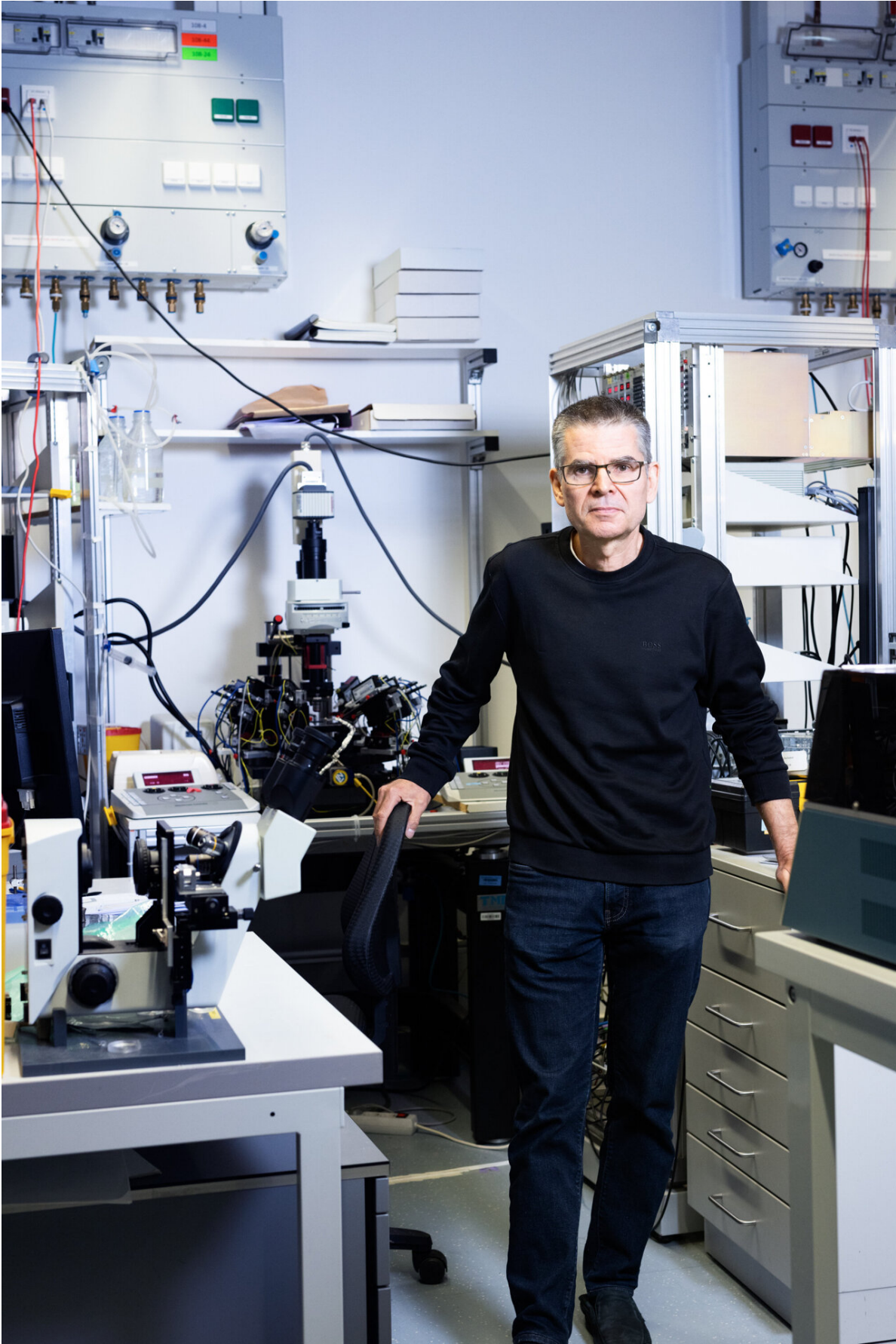
500,000 neurons in one model

In previous research, Jonas and his team measured crucial parameters of synapses, the connecting points of [neurons](#), and connectivity rules, which are required for understanding information processing in the

hippocampal network in mice and rats. Now, Jonas used these real-live parameters to build an accurate [model](#) of the network—a challenge, as usually, models of brain circuits are built with just 10 to 1,000 neurons. The dentate gyrus of rats, however, contains 500,000 excitatory neurons, called granule cells. "In scaled-down models, we cannot plug in the measured synaptic parameters of the neuronal circuit. But as we wanted to mimic what happens in the brain and use synaptic measurements we obtained previously, we implemented a network in its full size, with 500,000 granule cells."

Inhibition separates patterns

Using this computer model, Jonas tested different hypotheses of how pattern separation works. "With this model we cannot just copy biology, but systematically change parameters and disentangle factors. This allows us to understand computations in the brain, and how biological factors support or limit computation."



Professor Peter Jonas uncovered a new mechanism of pattern separation. Credit: © Peter Rigaud/IST Austria

Historically, based on data from the cerebellum, neuroscientists thought pattern separation is based on expansion: A pattern is projected from a smaller number of neurons to a huge number of neurons in the next layer of processing. This would expand the pattern and make it easier to spot differences. While expansion is a potential mechanism in the cerebellum, it is less likely to be at work in the hippocampus, where [granule cells](#) converge on a smaller number of CA3 neurons in the next layer.

"Clearly, expansion can't be the only mechanism in the hippocampus," Jonas states. "Evidence from our realistic model suggests that inhibition—active neurons stopping other neurons from firing—plays an important role." Mathematically, it has been shown that reducing activity in a network makes it easier to distinguish differences between patterns. Using the hippocampus model, Jonas interrogated the role of inhibition. "When inhibition is part of the model, patterns are robustly separated. But when we take inhibition out, this is no longer the case. This modeling data changes the historical view from code expansion to a mechanism based on inhibition."

The new data also explains a result from previous research which has left Jonas puzzled. "Previously, we found that inhibition in the dentate gyrus is locally constrained. Activated neurons only inhibit other cells within a 300-micrometer radius. We have long asked what might be the functional role of such focal inhibition." The network model shows that such focal inhibition can better separate patterns than global inhibition,

in which the entire network's activity is dampened down. Speed is essential in pattern separation, and focal inhibition reduces delays: Neurons in a pattern switch on and very rapidly inhibit surrounding cells, ensuring that other patterns are not turned on. "It is a cool solution, but not very intuitive, and we could only figure this out using the model," Jonas points out. As a next step, Jonas plans to transfer results back to the biological system and carry out behavioral experiments, further exploring how [inhibition](#) contributes to [pattern separation](#).

More information: Peter Jonas, How connectivity rules and synaptic properties shape the efficacy of pattern separation in the entorhinal cortex–dentate gyrus–CA3 network, *Nature Computational Science* (2021). DOI: [10.1038/s43588-021-00157-1](https://doi.org/10.1038/s43588-021-00157-1).
www.nature.com/articles/s43588-021-00157-1

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