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The scientists responsible for key breakthroughs in optogenetics distinguished with the Frontiers of Knowledge

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### award<sub>ence</sub>

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Communications

26 January 2016 Neuroscientists Edward Boyden, Karl Deisseroth and Gero Miesenböck have been distinguished with the BBVA Foundation Frontiers of Knowledge award in Riomedicine

(http://www.fbbva.es/TLFU/tlfu/ing/home/index.jsp) for their groundbreaking work in optogenetics, a technique that uses light to activate or deactivate proteins in neurons, enabling control over their activity with an unprecedented level of accuracy.

In just five years, thousands of groups across the globe have started using optogenetics to research brain functions such as sleep, appetite, decision making, the perception of time or the formation of memories, as well as to understand the mechanisms that trigger illnesses such as epilepsy, Parkinson's disease, depressions and even certain forms of blindness.

The beauty of optogenetics is that it allows the selective control of neural activity simply by applying light of the right wavelength. Before, the most widely used methods to study the living brain could modulate the activity of hundreds of thousands of neurons, but with little selectivity. With optogenetics, it is possible to act exclusively on neurons treated previously with light-sensitive proteins, according to the behavior being tested.

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<u>inifo/1KBTFA2&text=Edward Boyden</u>)
<u>Edward Boyden</u> (Plano, Texas, United States, 1979) completed his BS studies in physics and electrical engineering and computer science at Massachusetts Institute of Technology (MIT) before obtaining a PhD in neuroscience at Stanford University, where he worked alongside Karl Deisseroth. In 1996, he joined the research staff at MIT, where he remains to this day as professor of Biological Engineering and Brain and Cognitive Sciences. He leads

the institutions synthetic neurobiology group, developing tools that enable the systematic mapping and repair of the brain and other complex biological systems, and is also an investigator at the MIT McGovern Institute for Brain Research and associate professor at the MIT Media Lab. He has been co-director since 2014 of the MIT Center for Neurobiological Engineering.

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(https://twitter.com/share?url=http://bbva.info/1KBTFA2&text=Karl Deisseroth)

Karl Deisseroth (Boston, Massachusetts, United States, 1971)



studied biochemical sciences at Harvard than went on to complete an MD at Stanford, where he also obtained his PhD in neuroscience. After finishing his PhD dissertation, he opted for a residency in psychiatry, and today continues to combine patient consultations with a devotion to basic research. He has spent most of his career at Stanford University, where he is currently DH Chen Professor of Bioengineering and Professor of Psychiatry and Behavioral Sciences. He served on the committee behind the BRAIN Initiative launched by the Obama administration in April 2013 to "accelerate the development and application of new technologies that will enable researchers to produce dynamic pictures of the brain that show how individual brain cells and complex neural circuits interact at the speed of thought."

(https://twitter.com/share?url=http://bbva.info/1KBTFA2&text=Gero\_Miesenböck)
Gero\_Miesenböck (Braunau, Austria, 1965) studied medicine at



the University of Innsbruck, where he also obtained his PhD.
During a three-month stay at the University of Umeå, Sweden
(1989). he came across the work of American biochemist James
E. Rothman (Nobel Prize in Medicine 2013). In 1992, he moved to
New York to work with Rothman at the Memorial Sloan-Kettering
Cancer Center. Following three years as an Associate Professor of
Cell Biology at Yale University, in 2007 he joined the faculty of the
University of Oxford, where he combines the post of Waynflete
Professor of Physiology with the leadership of the Centre for
Neural Circuits and Behaviour (CNBC), which he also founded.

## A revolutionary technique that has found its way around the world

Ed Boyden, (Plano, Texas, United States, 1979), a professor at Massachusetts Institute of Technology, used the following simile in conversation yesterday, after hearing of the award: "If we imagine the brain as a computer, optogenetics is a key that allows us to send extremely precise commands. It is a tool whereby we can manipulate the brain with exquisite precision."

In 2002, Miesenböck was the first to show that neural activity could be modulated by light. He began with cultured cells, but realized immediately that "this was a technology with immense transformative power." Miesenböck recalls that **the reviewers of his first submission "completely missed the point"**; something he ascribes to the fact that "innovative techniques always take time to filter through."

His second "breakthrough" moment was in 2005, when he and postdoctoral assistant Susana Lima managed to employ the technique in a living organism, concretely a fruit fly. By activating just two of its hundreds of thousands of neurons, **optogenetics** triggered the response that made the insect fly away.

But Miesenböck's technique had major drawbacks: the proteins used caused only modest neural activation, and the process was unlikely to be suitable for large-scale application.

The key to a solution lay with Karl Deisseroth and Ed Boyden. Like Miesenböck, they tried working with light-sensitive proteins, but without ever cracking the problem. The two men returned to the fray in 2004, only this time using a protein recently isolated in a species of green alga, with far superior light responsiveness to those employed by Miesenböck.

Deisseroth and Boyden's paper appeared in 2005. Since then, the technique has undergone successive refinements, with **proteins**, for instance, that respond at different speeds or to different types of lights, extending the range of brain functions that can be studied.

Boyden recounts how after the 2005 paper, rejected in turn by *Science* and *Nature*, he was turned down for jobs by at least half of the institutions he applied to: "At the time neuroscientists were distrustful of neurotechnology."

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LAST UPDATE

02 February 2016 Finnish ecologist Ilkka Hanski has been distinguished with the BBVA Foundation Frontiers of Knowledge Award in Ecology and Conservation Biology

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(http://www.fbbva.es/TLFU/tlfu/ing/microsites/premios/fronter**its**/index.jsp) for opening up a new branch of ecology - g+ metapopulation biology - which combines field work and mathematical modeling to predict how species will be affected by the pressing problem of habitat fragmentation by human action.

Hanski has developed mathematical models that allow predicting the viability of populations of many species and habitats. In its statement, the award's international jury emphasizes that "the impact of Professor Hanski's work increases as habitats become more and more fragmented by anthropogenic influence."

"Human activity increases the degree of fragmentation, so it is vital to know what the consequences are. **Metapopulation theory helps us devise strategies to improve population survival.** A typical example would be the optimal design of a reserve, deciding how it should be organized so it can fulfill its function with the least possible impact on species," explains Hanski.

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His contributions inform many of the conservation initiatives around us today, from the design of protected areas to the creation of biological corridors or environmental impact studies prior to major infrastructure developments. It is also used to define conservation strategies in fragmented habitats or habitats affected by forces like deforestation, urbanization or climate change.

Mathematical models of metapopulation biology allow us to quantify the degree of habitat fragmentation a given species can withstand. In other words, to "determine the critical threshold beyond which fragmentation is fatal and the species is lost," as Hanski himself explains in this video.

Veerabhadran Ramanathan, Frontiers of Knowledge award in Climate Change (https://info.bbva.com/en/n ramanathanfrontiers-knowledgeaward-climatechange/)

## Human health applications: Cancer, epidemics, and immune system

As Hanski explains, "the metapopulation concept, has been incorporated into many different areas of research." In cancer research, for instance, it is used to study the behavior of tumor cell populations. In epidemiology, knowing the critical size of a metapopulation network can help fight the spread of a disease.

Hanski himself has studied the link between the human immune system and the various microorganism communities that populate the human body (microbiome). According to his research, the biodiversity degree to which individuals are exposed and the behavior of their microbiome have implications for the state of their immune system. "After all, for our microbiome we ourselves are fragmented habitats, and it is intellectually fascinating to work with colleagues from other fields, such as immunology."

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**y** (https://twitter.com/share?url=http://bbva.info/1PwhkHs&text=Butterflies and beetles, a source of inspiration)



Hanski (Finland, 1953) first got interested in population distribution in the late 1970s, as a doctorate student at the University of Oxford (United Kingdom). His attention was drawn to dung beetles, and the way different species clumped together in cattle pats. He began using mathematical models to describe the behavior of each species.



At the end of the 1980s, back home in Finland, he commenced studying the Glanville fritillary butterfly (Melitaea cinxia). He was inspired in this endeavor by the visit to Finland of Stanford entomologist Paul Ehrlich (United States), a Frontiers of Knowledge laureate

(http://www.fbbva.es/TLFU/tlfu/ing/microsites/premios/fronteras/galardonados/2013/ecologia.jsp),although Hanski himself had been a keen butterfly collector in his

He selected a large study area in the Åland Islands, in the Baltic Sea, where thousands of dry meadow patches provided the ideal fragmented habitat. From the early 1990s, Hanski and his students conducted an annual census of the butterflies present in each meadow, to refine their models and test their predictions.

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