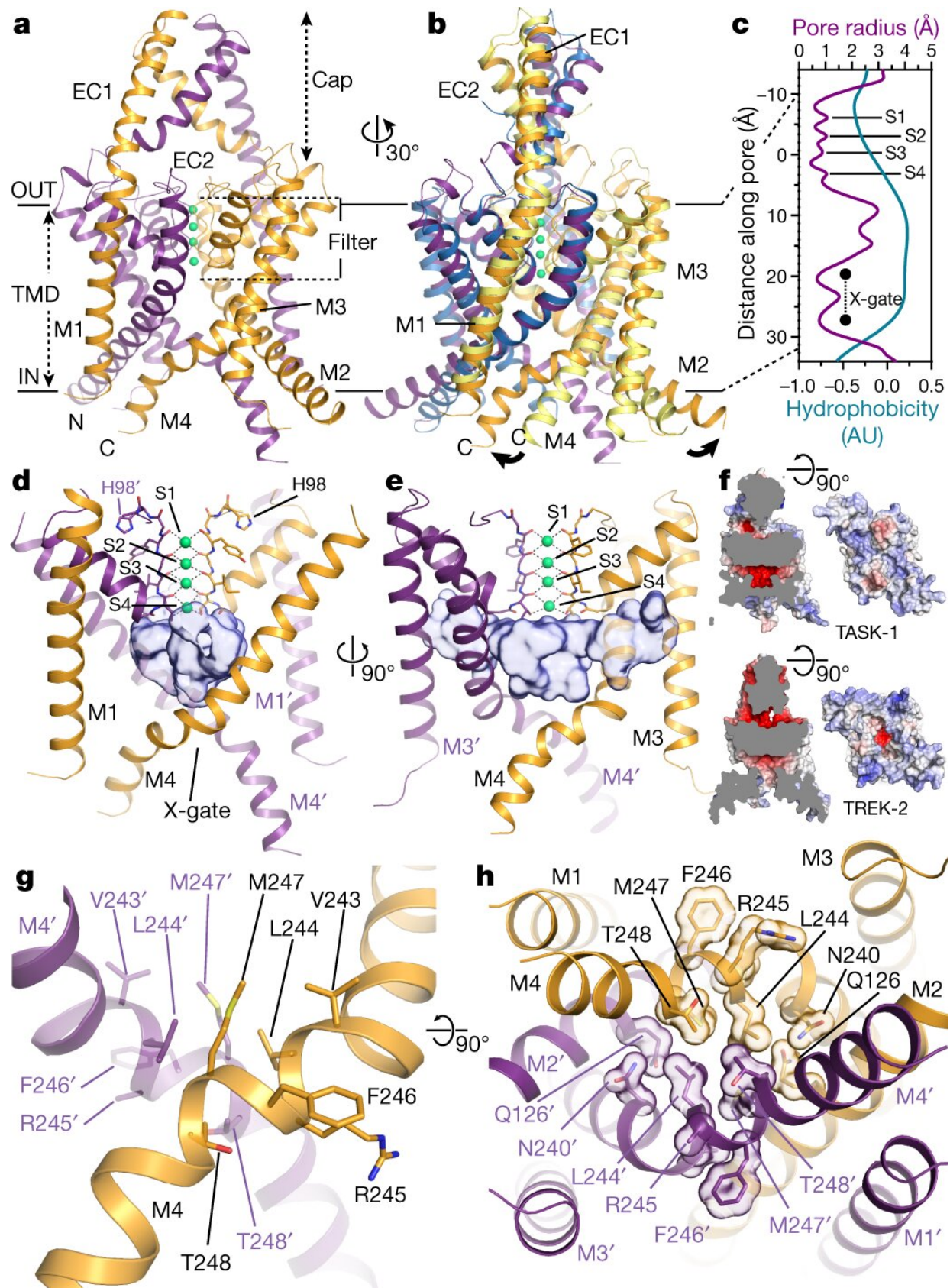


Three-dimensional molecular structure of TASK channel described

May 1 2020, by Bob Yirka



TASK-1 has a unique crossed-helix X-gate. a, Cartoon showing the structure of TASK-1, with chain A in gold, chain B in purple and potassium ions in green, viewed from the membrane. b, Superposition of the structures of TASK-1 (gold and purple) and TREK-2 (down state (PDB: 4XDJ)) (yellow and blue). c, Radius and hydrophobicity of the TASK-1 pore. AU, arbitrary units. d, e, The X-gate enclosing the vestibule and fenestrations (depicted with semi-transparent surfaces), viewed from the membrane plane (d) and rotated 90° (e). f, Electrostatic surfaces of TASK-1 and TREK-2 (PDB: 4XDJ) shown as a cross-section perpendicular to the membrane and seen from below. g, h, The X-gate viewed from the plane of the membrane (g) and below the membrane (h). Credit: *Nature* (2020). DOI: 10.1038/s41586-020-2250-8

A team of researchers from the University of Oxford, the University of Marburg and Bayer Pharmaceuticals has developed a way to describe the three-dimensional molecular structure of TASK channels. In their paper published in the journal *Nature*, the group describe the technique they used to describe the potassium channel in detail for the first time and what they discovered as they were doing so.

TWIK-related acid-sensitive potassium (TASK) channels are members of the general [potassium](#) channel family—what sets them apart are their two-pore structure and their location—they are found in neurons, smooth vascular muscle cells and cardiomyocytes (heart muscle cells). Potassium channels are portals through cell membranes that allow or bar the transport of ions. They do so through the use of gates that physically open or close. Such channels are found in almost all [living organisms](#)—they help regulate electrical signals and nerve activity. In this new effort, the researchers sought to learn more about the structure of TASK channels, most specifically, why they are able to bind inhibitors (chemicals that urge the channels to bar the passage of ions) so well. Medical scientists have developed such inhibitors to treat some forms of

respiratory failure, cardiac fibrillation and some sleep disorders such as apnea.

To learn more about the structure of TASK channels, the researchers studied them using X-ray crystallography—it allowed them to get an atom-to-atom view of the ion portals—and to find something unexpected. In most [potassium channels](#), there is a gate with four helices in parallel located below a selectivity filter, but to date, no such lower gates had been seen in TASK channels. In this new effort, the researchers found evidence of just such a lower gate, which they named: X-gate. And it was created by interactions between two transmembrane helices at the entrance to the vestibule.

Study of the [X-gate](#) showed it kept inhibitors in the entrance part of the [channel](#) and that explained why inhibitors have been so notoriously difficult to wash out. They note that the discovery of the X-gate's existence also helps to explain other aspects of TASK channels and will likely help with the development of new drugs to treat lung and sleep disorders.

More information: Karin E. J. Rödström et al. A lower X-gate in TASK channels traps inhibitors within the vestibule, *Nature* (2020). [DOI: 10.1038/s41586-020-2250-8](https://doi.org/10.1038/s41586-020-2250-8)

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Citation: Three-dimensional molecular structure of TASK channel described (2020, May 1) retrieved 20 April 2024 from <https://phys.org/news/2020-05-three-dimensional-molecular-task-channel.html>

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