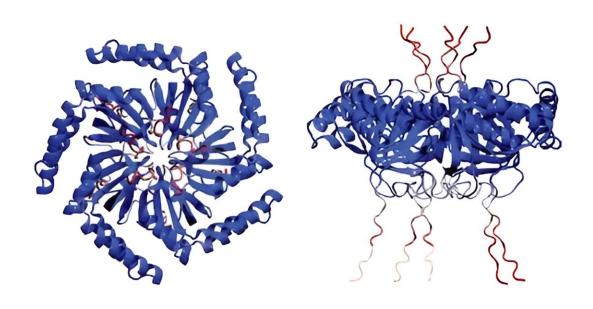


Study identifies protein responsible for gas vesicle clustering in bacteria

March 29 2024, by Patrick Kurp



Simulated configuration of the protein structure. Credit: Rice University

Gas vesicles are hollow structures made of protein found in the cells of certain microorganisms, and researchers at Rice University believe they can be programmed for use in biomedical applications.

"Inside cells, gas vesicles are packed in a beautiful honeycomb pattern. How this pattern is formed has never been thoroughly understood. We



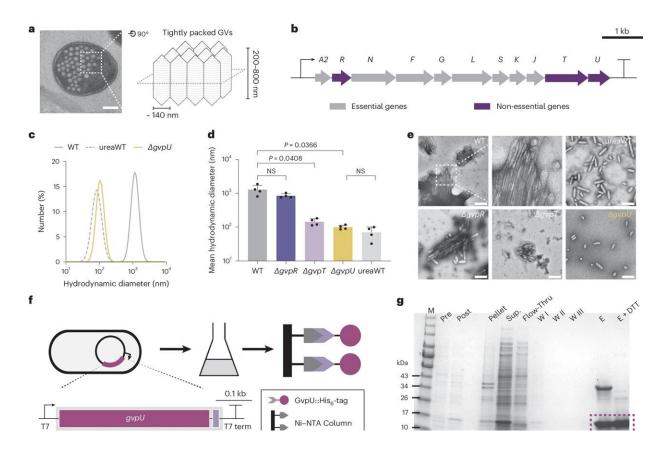
are presenting the first identification of a <u>protein</u> that can regulate this patterning, and we believe this will be a milestone in molecular microbiology," said George Lu, assistant professor of bioengineering and a Cancer Prevention and Research Institute of Texas scholar.

Lu and colleagues have published their findings in <u>a paper</u> published in *Nature Microbiology*. The lead author is Zongru Li, a fourth-year bioengineering doctoral student in Lu's Laboratory for Synthetic Macromolecular Assemblies.

"Gas vesicles are cylindrical tubes closed by conical end caps," Li said.
"They provide buoyancy within the cells of their native hosts."

The vesicles are found naturally in five phyla of bacteria and two groups of archaea (single-cell organisms). Most are restricted to planktonic microorganisms often found in freshwater ponds. The recent engineering of vesicles has led to several applications, including reporter gene imaging, acoustic control, and payload delivery.





Gene knockout screening determines GvpU is essential for the clustering of GVs, and purified GvpU reconstitutes the GV cluster. Credit: *Nature Microbiology* (2024). DOI: 10.1038/s41564-024-01648-3

Co-author Yifan Dai, assistant professor of biomedical engineering at McKelvey School of Engineering at Washington University in Saint Louis, said they were drawn to the research with the question of why the vesicles can form in the honeycomb pattern.

With help from his WashU colleague Alex Holehouse and colleagues from Duke University, Ashutosh Chilkoti, and Lingchong You, the team of researchers found that this pattern is the most efficient use of space and the cluster form plays a part in how it functions.



Most notably, these protein clusters formed in subsaturated solution, a previously identified new form of biological structure, and that drives the organization of these vesicles. The bottom line is that they found the function behind this mysterious new form.

"These teams led by Lu lab found that a unique form of protein clusters exclusively assembled in subsaturated solution drives the clustering behaviors," said Dai. This adds to the line of evidence on how phase transition affects cellular organization and cellular functions, he added.

Lu and his team, using genetic, biochemical, and imaging approaches, are exploring protein nanostructures. Gas vesicles stabilize the air bubbles inside the bacterial cytosol—the fluids inside the cells—and provide a liquid-gas interface that can be used for ultrasound or MRI contrast.

"In our lab, we are leveraging the power of synthetic biology to expand the applications of these protein nanostructures," Li said. "By engineering genes and cells, we aim to build gas vesicles that perform even more efficiently in biotechnological and <u>biomedical applications</u>."

More information: Zongru Li et al, Phase transition of GvpU regulates gas vesicle clustering in bacteria, *Nature Microbiology* (2024). DOI: 10.1038/s41564-024-01648-3

Provided by Rice University

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