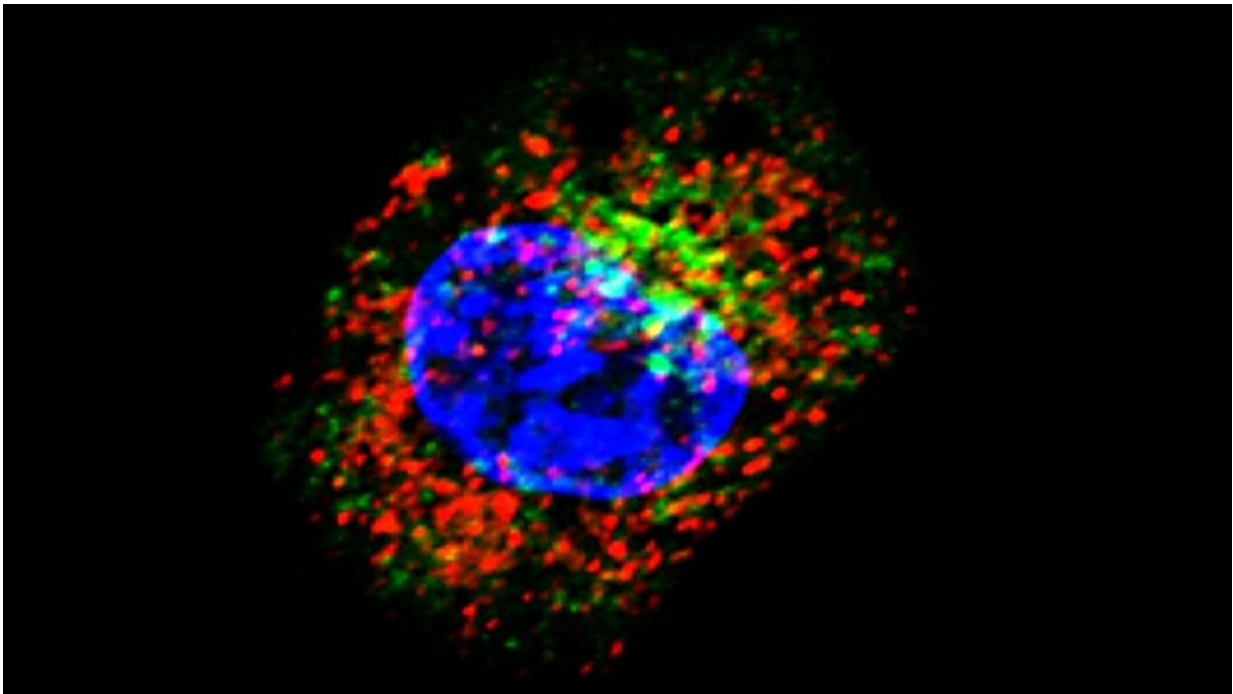


# How a little-known glycoprotein blocks a cancer cell's immune response

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The interaction between STC1 and calreticulin. Credit: University of Michigan Rogel Cancer Center

It was an unexpected discovery that started with an analysis of more than 1,000 genes. The question: why game-changing cancer immunotherapy treatments work for only a fraction of patients.

The analysis shone a light on one that popped up repeatedly in patients

and mouse models that did not respond to immune checkpoint therapy: stanniocalcin-1, a glycoprotein whose role in both tumors and immunology is largely unknown.

By following the trail from this surprising thread, a University of Michigan Rogel Cancer team uncovered how stanniocalcin-1, or STC1, works inside the cell to block a cellular "eat-me" signal that typically triggers the [immune system](#) to produce T [cells](#) to fight the [tumor](#). The findings, published in *Cancer Cell*, provide a potential target to improve immune responses to cancer.

"We believe STC1 is a checkpoint inside the cell. It's an eat-me blocker—it blocks macrophages and [dendritic cells](#) to eat dying or dead [cancer cells](#). We think that if we can target the STC1 pathway, it would release the blocked eat-me signal," says study senior author Weiping Zou, M.D., Ph.D., Charles B. de Nancrede Professor of Pathology, Immunology, Biology, and Surgery at the University of Michigan.

Zou and colleagues were drawn to STC1 in part because so little was known about its role in cancer. This provided a potentially interesting opportunity, but also some difficulty as they had to start at the very beginning to understand whether STC1 was causing the poor [immune response](#) or whether it was just a bystander.

They embarked on a lengthy process, first showing that STC1 was linked with low activation of T cells and worse survival in melanoma patients treated with immunotherapy. They checked against the Cancer Genome Atlas database and found high levels of STC1 tied to worse survival in 10 different [cancer](#) types. The finding also panned out in mouse models.

From there, the researchers used mouse models to show that STC1 within tumors was dampening the anti-tumor T cell response by impairing the antigen presenting cells, which are essential for triggering

T cells. They showed that tumor STC1 was stopping the process of macrophages, a type of antigen presenting cell, from eating dying tumor cells—a process key to presenting antigen to T cells and activating them.

Specifically, the tumor STC1 traps a key eat-me signal called calreticulin, or CRT. Without sufficient surface CRT, the macrophages won't efficiently eat the dead tumor cells. Unblock the CRT and it unblocks this process. This suggests that targeting the interaction between STC1 and CRT might be a path toward making immunotherapy more effective.

It's an unusual mechanism. Most immune checkpoint therapies are based on direct surface interactions with T cells.

"What we are talking about is before anything has happened. Before the T cells were activated, the tumors have already implemented strategies so they cannot be captured. This may be why some patients are resistant to immunotherapy: their tumors express too much STC1. When you block the eat-me signal, the antigen presenting cells cannot do their job," Zou says.

Targeting the STC1 and CRT interaction inside the cell is trickier than if it were on the cell surface. It means a typical antibody approach will not work. Instead, Zou and colleagues are investigating whether they can develop a small compound that would penetrate the cell and interfere with the STC1-CRT interaction.

**More information:** *Cancer Cell* (2021). [DOI: 10.1016/j.ccell.2020.12.023](https://doi.org/10.1016/j.ccell.2020.12.023)

Provided by University of Michigan

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