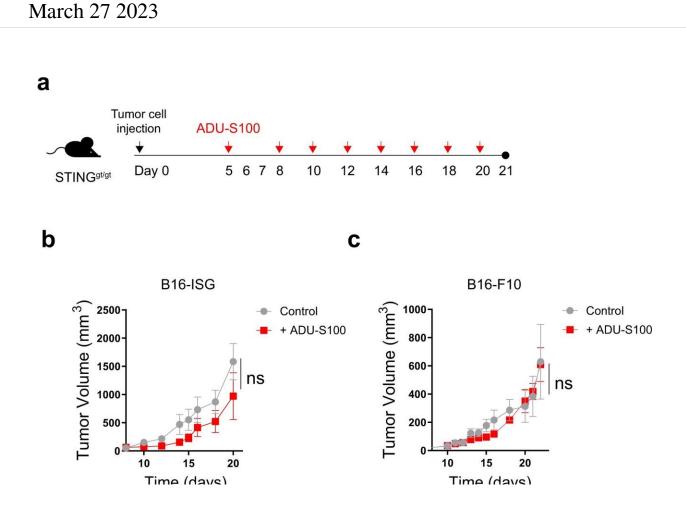


Researchers discover two-pronged approach to stimulate STING antitumor activity



Schematic of the STING agonist treatment schedule. Groups of STING^{gt/gt} mice were injected subcutaneously with 1.5×10^5 B16-ISG or B16-F10 on day 0. On days 5, 8, 10, 12, 14, 16, 18 and 20 following tumor injection, tumor-bearing mice were intratumorally treated with either PBS or 50 µg ADU-S100 (a). Tumor growth curves of B16-ISG (b) and B16-F10 (c) in STING^{gt/gt} mice treated with PBS control or ADU-S100 as indicated in (a). Data are shown as the mean \pm SEM and are representative of two independent experiments. n = 4 and 5 mice in (b) and n = 5 and 6 mice in (c) for ADU-S100 and Control groups,



respectively. The frequency of CD8⁺ cells within the CD45⁺ population in B16-ISG (d) and B16-F10 (f) tumors treated with PBS control or ADU-S100 (n = 4 mice per group). Data are shown as mean \pm SD and are representative of two independent experiments. Statistical significance was determined by a two-sided t test (ns, not significant). Representative flow cytometry plots for d and f are shown in e and g, respectively. Credit: *Nature Communications* (2023). DOI: 10.1038/s41467-023-37217-1

Immunotherapies have greatly improved the outcomes of many patients with melanoma. But there is still a need for new approaches for the subset of patients who do not respond well to this type of therapy. Moffitt Cancer Center researchers are looking at new targets to help inhibit tumor development and promote antitumor immunity, one being the STING signaling pathway. In a new article published in *Nature Communications*, a team of Moffitt and University of Miami Miller School of Medicine investigators demonstrate that targeting the STING pathway with a combination strategy results in improved antitumor activity.

The STING pathway is a key regulator of immune responses to viruses and bacteria and contributes to antitumor immunity. Inhibition of STING signaling is observed in several <u>cancer</u> cell lines, and expression of STING protein decreases as some cancer types progress, such as melanoma. These observations suggest that agonist drugs, or drugs that activate STING signaling within the tumor environment, may have antitumor effects.

"This hypothesis has been confirmed in many pre-clinical laboratory studies. However, STING agonists have had disappointing results in human clinical trials, suggesting that an unknown mechanism is contributing to these poor responses," said lead study author Rana Falahat, Ph.D., a research scientist in the Immuno-Oncology Program at



Moffitt. "This is what prompted our study."

The Moffitt and University of Miami team wanted to improve their understanding of STING signaling and investigate how to improve the antitumor activity of STING agonists. They previously demonstrated that the DNA region that controls STING protein expression is modified by a process called methylation, during which chemical <u>methyl</u> groups are added to the DNA. This modification reduces the protein level of STING, resulting in less STING protein to inhibit <u>tumor development</u>. The researchers hypothesized that drugs that block methylation and increase STING expression levels in tumor cells may work in conjunction with STING agonists to kill <u>cancer cells</u>.

The researchers performed experiments in mouse models of melanoma with defective STING signaling either in the tumor cells or the surrounding immune cells. Similar to the human STING gene, they discovered that the mouse STING gene was also regulated by methylation. When the researchers treated cells with drugs that blocked methylation, STING expression and signaling increased, leading to the production of chemical messengers that enhanced antitumor immunity. They also demonstrated that treatment of mice with methylation inhibitors and STING agonists resulted in improved antitumor activity that was dependent on the presence and activity of immune cells called CD8 T cells.

These observations reveal that reactivating STING signaling in the <u>tumor</u> <u>cells</u> with methylation inhibitors can shape the therapeutic response to STING agonists. The researchers hope these preclinical findings will lead to better treatments for cancer patients in the clinic.

"Although additional work will be necessary to further identify optimal dosing levels and therapeutic schedules of each component, insights from our study provide a framework to design proper clinical treatment



modalities with appropriate patient selection in melanoma and perhaps other <u>solid tumors</u>," said James J. Mulé, IPh.D., senior author, associate center director for Translational Science and the Michael McGillicuddy Endowed Chair for Melanoma Research and Treatment at Moffitt.

More information: Rana Falahat et al, Epigenetic state determines the in vivo efficacy of STING agonist therapy, *Nature Communications* (2023). DOI: 10.1038/s41467-023-37217-1

Provided by H. Lee Moffitt Cancer Center & Research Institute

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