

Scientists uncover a new approach for treating aggressive cancer by inhibiting chromatin-modulatory enzyme EZH2

24 February 2022



Credit: CC0 Public Domain

Researchers at the University of North Carolina at Chapel Hill and the UNC Lineberger Comprehensive Cancer Center have uncovered a new role of a chromatin-modulatory enzyme, termed EZH2, during cancer development. They then developed a new therapeutic approach with a potent small-molecule inhibitor of this enzyme.

Certain subtypes of blood cancers such as acute leukemias rely on multiple mechanisms for sustaining growth of aggressive [cancer](#) cells. Notably, these mechanisms include those driven by EZH2, a chromatin-modulatory enzyme, and cMyc, a prominent cancer-causing factor. UNC researchers now show that these two factors can directly associate with one another, modulating cancer-cell-specific programs of gene expression.

To develop pharmacological means of targeting both EZH2 and cMyc, they teamed with the chemical biologists at Icahn School of Medicine at Mount Sinai and designed a new small-molecule,

MS177, based on the proteolysis-targeting chimera (PROTAC) technology. MS177 targets both EZH2 and cMyc and thus inhibit cancer growth.

Their findings are published online in *Nature Cell Biology*.

"EZH2 plays a very important role during cancer progression and is a known target suitable for drug development," said UNC Lineberger's Greg Wang, Ph.D., associate professor of Biochemistry and Biophysics and Pharmacology at the UNC School of Medicine and co-lead author of this research article. "We are amazed by the efficiency of small-molecule PROTAC in simultaneously targeting EZH2 and cMyc in cancer cells."

They found that EZH2 possesses two different binding patterns on chromatin in acute leukemia cells, eliciting two distinct gene-regulatory programs (Figure 1). On the one hand, EZH2 forms a canonical protein complex termed PRC2, leading to gene repression at a set of genomic regions; on the other hand, EZH2 interacts with cMyc to activate gene expression at genomic sites distinctive from the above ones. "This explains why the current small-molecule inhibitors of EZH2 cannot block EZH2 completely. PROTAC addresses this gap," said Jun Wang, Ph.D., postdoctoral researcher at UNC Lineberger and co-first author of the work.

MS177 achieves on-target effect in cancer cells and exhibits profound tumor killing effects, the researchers report. "Compared to the existing enzymatic inhibitors, MS177 is more likely to behave much better for the treatment of patients with acute leukemias. To our knowledge, an agent for dual targeting of EZH2 and cMyc has not been developed before. cMyc is hard to 'drug,'" Greg Wang said. "MS177 thus represents a promising candidate for treating other cancers depending on

the above tumorigenic pathways."

More information: Jun Wang et al, EZH2 noncanonically binds cMyc and p300 through a cryptic transactivation domain to mediate gene activation and promote oncogenesis, *Nature Cell Biology* (2022). [DOI: 10.1038/s41556-022-00850-x](https://doi.org/10.1038/s41556-022-00850-x)

Provided by UNC Lineberger Comprehensive Cancer Center

APA citation: Scientists uncover a new approach for treating aggressive cancer by inhibiting chromatin-modulatory enzyme EZH2 (2022, February 24) retrieved 9 July 2022 from <https://medicalxpress.com/news/2022-02-scientists-uncover-approach-aggressive-cancer.html>

This document is subject to copyright. Apart from any fair dealing for the purpose of private study or research, no part may be reproduced without the written permission. The content is provided for information purposes only.