

Pediatric cancers share stalled gene-managing enzyme

October 16 2020, by Chris Barncard



Credit: Unsplash/CC0 Public Domain

A wildly out-of-place protein leads to haywire cells in a particularly troublesome type of rare early childhood cancer, according to University of Wisconsin–Madison researchers.

Found in the base of the brain, posterior fossa type A ependymomas

tumors are difficult to remove via surgery and prove fatal in more than a quarter of children within five years of diagnosis. A study published this week in the journal *Molecular Cell* describing the role in the brain tumor cells of a wayward protein called EZHIP, which belongs in ovaries and testes, may suggest a target for new treatments.

Peter Lewis, a UW–Madison professor of biomolecular chemistry who led the research, studies chromatin dynamics: the [protein molecules](#) and processes that pack and unpack six feet of genomic DNA in the nucleus of every human cell. In recent years, his research group and collaborators have shown that specific mutations changing one group of packaging proteins called histones—the molecular spools around which DNA winds in the nucleus— can change the way a cell's genes are expressed to make the cell cancerous.

"These mutant histones found in human cancers—we call them oncohistones— make up a tiny proportion of the total histone content of tumor cells," Lewis says. "But these oncohistones really punch above their weight because of how they work."

Histones normally help manage gene expression by unraveling parts of DNA strands only under appropriate biochemical circumstances.

"Many parts of the genome remain tightly compacted throughout most of the lifetime of an animal," Lewis says. "They might only become de-compacted, to allow DNA in that part of the genome to be used by the cell's machinery, during a very transient window of development in one specific cell type. This helps regulate genomic DNA access. And that, in turn, affects gene expression. And that, in turn, establishes and maintains cell identity."

A mutated histone, however, may reveal or conceal genes at inappropriate times.

"When things go awry—for example, a specific mutation in histone proteins or mutations to the machinery that compact and de-compact the chromatin— this can have dire consequences for the cell," Lewis says.

In the case of a lethal brainstem tumor called diffuse intrinsic pontine glioma, or DIPG, Lewis and others discovered in 2013 that an oncohistone called K27M inhibits the activity of an enzyme called PRC2. Disruptions to PRC2, which itself helps control the appropriate expression of genes, can cause cells to turn cancerous.

In their new study, the researchers—including UW–Madison doctoral student Siddhant Jain, biomolecular chemistry professor Melissa Harrison, and collaborators from McGill University Health Center—found that the molecules driving DIPG tumors and PFA ependymomas function in a remarkably similar manner.

"The PFA ependymomas look, molecularly, just like DIPG. They had all the tell-tale signs," Lewis says. "We looked and looked and looked, but PFA ependymomas didn't have the same K27M oncohistone. The cells that give rise to this tumor have, instead, somehow turned on EZHIP."

That shouldn't happen. EZHIP is a protein that's normally turned on only in cells of the ovaries and testes—nowhere near brain tumors.

"It's not supposed to be present in the brain, and we don't understand how it's turned on," Lewis says. "But we now know what it does when it is present. We found that it acts just like the K27M oncohistones. It inhibits the same PRC2 complex."

When PRC2 is active, it is both reading chemical modifications from histones and writing new ones to histones to promote proper chromatin compaction and cell function.

"EZHIP and K27M cause writer's block (in PRC2)," Lewis says. "And it's not just any PRC2 which is inhibited, it's a specific activated form—a super writer—that's blocked."

Zeroing in on PRC2 and its effects may present new ways to treat these cancers.

"Because we understand how a form of PRC2 is inhibited by these tumor-driver proteins, we can now develop a screen for small molecules that target this specific form of the enzyme," Lewis says. "EZHIP's very specific normal location in the gonads and K27M's presence only in tumor [cells](#) also make them potential candidates for anti-tumor vaccines."

In a second study, published this week in the *Proceedings of the National Academy of Sciences*, the same group of researchers discovered that PRC2 activity is affected by a different oncohistone called G34 in 92 percent of giant-cell tumors of the bone, another pediatric [tumor](#). In these tumors, the researchers found that G34 oncohistones allow PRC2 to be overactive in parts of the genome where it shouldn't normally function.

"These specific bone tumors are generally less deadly than DIPG, but understanding the relevance of PRC2 in their tumorigenesis presents an opportunity to explore alternative therapies," says Lewis.

More information: Siddhant U. Jain et al. H3 K27M and EZHIP Impede H3K27-Methylation Spreading by Inhibiting Allosterically Stimulated PRC2, *Molecular Cell* (2020). [DOI: 10.1016/j.molcel.2020.09.028](https://doi.org/10.1016/j.molcel.2020.09.028)

Provided by University of Wisconsin-Madison

Citation: Pediatric cancers share stalled gene-managing enzyme (2020, October 16) retrieved 6 July 2024 from <https://phys.org/news/2020-10-pediatric-cancers-stalled-gene-managing-enzyme.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.