

Using Light to Fight Cancer: A Promising New Treatment Strategy

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Cancer cells often hijack normal cellular pathways to promote uncontrolled growth, invade surrounding tissues, and spread to distant parts of the body. A recent study has identified a protein called Aspartate γ -Hydroxylase (ASPH) as a potential target for future cancer therapies, offering new insights into how tumors grow and how they may be stopped.

What Is ASPH?

ASPH is an enzyme that is normally expressed at low levels in most adult tissues. However, many cancers, including breast, cervical, liver, and head and neck cancers, produce unusually high amounts of ASPH. Previous research has suggested that ASPH helps tumors become more aggressive by activating signaling pathways involved in cell growth, migration, and invasion. Because of its elevated expression in many tumor types and limited presence in healthy tissues, ASPH has emerged as an attractive target for cancer therapy.

Testing New ASPH Inhibitors

In the new study, researchers examined the effects of two experimental ASPH inhibitors, known as MO-I-1151 and MO-I-1182, in multiple human cancer cell lines derived from cervical, pharyngeal, and breast tumors. The goal was to determine whether blocking ASPH could slow cancer progression. The results were encouraging. Treatment with ASPH inhibitors significantly reduced:

- * Cancer cell proliferation (growth)
- * Cell migration
- * Cell invasion into surrounding tissues

These findings suggest that ASPH plays an important role in multiple aspects of tumor progression and that inhibiting its activity may help suppress aggressive cancer behavior.

Disrupting Key Cancer Signaling Pathways

The researchers also investigated how ASPH inhibition affected cellular signaling. They found that blocking ASPH disrupted both the canonical and non-canonical Notch1 signaling pathways, which are known to regulate cell growth, survival, and differentiation. The study also identified interactions with AKT signaling, another pathway frequently activated in cancer. These signaling changes were accompanied by a decrease in Cyclin D1, a protein that helps drive cells through the cell cycle and promotes proliferation. Reduced Cyclin D1 levels were

associated with an accumulation of cells in the G0/G1 phase of the cell cycle, preventing progression into DNA synthesis and cell division. In simple terms, inhibiting ASPH appeared to place the cancer cells into a growth arrest state, limiting their ability to multiply.

Not All Tumors Respond the Same Way

One of the most important findings from the study was that different cancer cell lines responded differently to ASPH inhibition. Some cell types showed stronger reductions in growth and invasion, while others displayed more modest effects. This variability highlights an important challenge in modern oncology: tumor heterogeneity. Even cancers that arise from the same tissue can possess different genetic and molecular characteristics, leading to different responses to treatment. The findings underscore the need for personalized approaches that match therapies to the specific biology of an individual's tumor.

Why This Matters

Targeted therapies have transformed cancer treatment by focusing on specific molecules that drive tumor growth. ASPH represents a promising candidate because it appears to influence multiple pathways involved in cancer progression while remaining relatively restricted in normal tissues.

Although these findings were generated in laboratory models and additional studies are needed before ASPH inhibitors can become standard treatments, the results provide further evidence that ASPH may serve as both a therapeutic target and a biomarker for aggressive cancers.

Looking Ahead

Future research will focus on determining which cancer types are most dependent on ASPH activity and whether ASPH inhibitors can improve outcomes in animal models and clinical trials. As scientists continue to unravel the complex networks that drive cancer growth, studies like this help identify new vulnerabilities that may one day lead to more effective and personalized cancer treatments.

Source

<https://pmc.ncbi.nlm.nih.gov/articles/PMC11134442/>

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