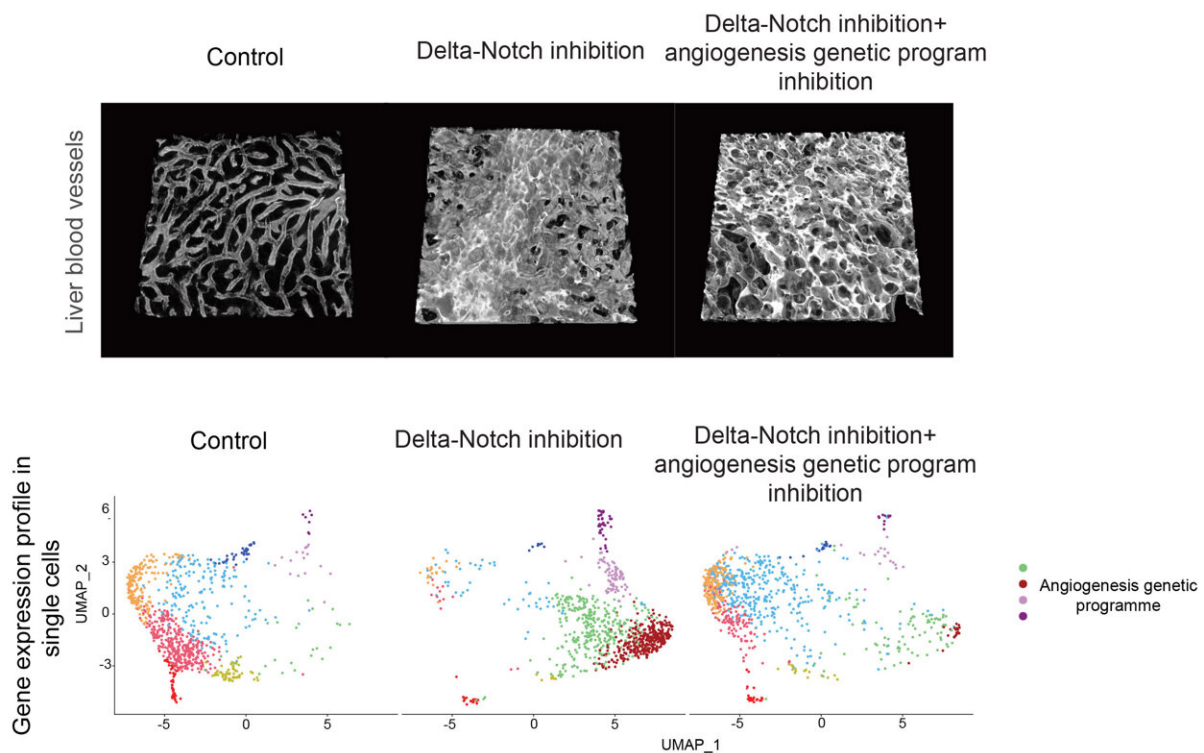


Scientists identify how some angiogenic drugs used to treat cancer and heart disease cause vascular disease

May 31 2023



Drugs that inhibit Delta-Notch signaling trigger significant alterations in vascular architecture and induce the expression of genes involved in angiogenesis. The changes in vascular architecture do not disappear when expression of the angiogenesis-related genes is blocked. Credit: CNIC

Research by scientists at the Centro Nacional de Investigaciones Cardiovasculares (CNIC) has demonstrated that the on-target molecular and cellular effects of medicines used to modulate the formation of new blood vessels (angiogenesis) in cardiovascular disorders and cancer are not responsible for the toxicity and vascular pathology triggered by these drugs. The study is published in *Nature Cardiovascular Research*.

Study leader Rui Benedito commented that, "Our results not only significantly increase our understanding of the biology of [blood vessels](#), but will also help in the selection of the most effective and safe way to modulate angiogenesis in ischemic tissues or in cancer."

The vascular system supplies oxygen and nutrients to the tissues and organs of the body. But the blood vessels do more than just conduct blood; they contribute actively to the physiology and homeostasis of all tissues and organs throughout life. Most blood vessels in the body are in an inactive state, which they maintain by expressing a large number of genes, including the genes of the signaling pathway mediated by Delta ligands and Notch receptors.

Several drugs have been developed in recent years that either block or induce angiogenesis in cardiovascular disorders or cancer.

A group of these compounds in clinical use inhibit different components of the Delta-Notch signaling pathway, which plays important roles in angiogenesis and in the maintenance of blood vessels in the inactive state. These compounds, by modulating the growth of blood vessels, efficiently block tumor growth. These compounds are also able to induce angiogenesis in ischemic tissues, thereby improving tissue regeneration and function.

However, these drugs can also cause [vascular injury](#) in organs with no previous disease, including the liver and heart, and this has reduced

clinical interest in their use.

Until now, this vascular toxicity was thought to be due to the expression of genes that promote angiogenesis, leading to the appearance of neoplasms or tumors in the affected blood vessels.

Thanks to the use of advanced genetic mouse models, high-resolution [confocal microscopy](#), and single-cell sequencing and proteomics techniques, the team discovered that the vascular toxicity linked to these drugs is instead due to a change in the vascular architecture that impedes correct blood flow.

"Our study shows that the vascular pathology that can result from treatment with these drugs is unrelated to the expression of genes involved in angiogenesis or the appearance of neoplasms," said Rui Benedito.

The researchers showed that these changes happen even when they blocked cell activation and the expression of angiogenesis-related genes.

Therefore, explained Rui Benedito, "although the neoplasms and the expression of genes associated with angiogenesis are associated with the change in vascular architecture, they are not the cause of this change."

First author Macarena Fernández Chacón explained that, "after analyzing different [genes](#) and drugs targeting blood vessels, we have found new ways to control pathological [angiogenesis](#) without significantly affecting vascular architecture in other organs, thus avoiding toxicity."

More information: Macarena Fernández-Chacón et al, Incongruence between transcriptional and vascular pathophysiological cell states, *Nature Cardiovascular Research* (2023). [DOI: 10.1038/s44161-023-00272-4](#)

Provided by Centro Nacional de Investigaciones Cardiovasculares Carlos III (F.S.P.)

Citation: Scientists identify how some angiogenic drugs used to treat cancer and heart disease cause vascular disease (2023, May 31) retrieved 21 June 2024 from <https://medicalxpress.com/news/2023-05-scientists-angiogenic-drugs-cancer-heart.html>

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