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Step by Step, Day by Day

Scientists uncover the series of molecular signals that guide earliest stages of artery development

BY ANNE D. HOLDEN, PHD

Not all circulatory systems are created equal. Vertebrates (animals with a backbone) have a so-called ‘closed’ circulatory system, made up of blood vessels. Many invertebrates (those with no backbone) have an ‘open’ system, in which blood and other fluids flows freely through the body. And some species, like flatworms, don’t have a circulatory system at all.

In fact, the ability to form blood vessels in the first place is one of evolution’s crowning achievements—and distinguishes vertebrates from the rest of the animal kingdom. The two types of blood vessels, arteries and veins, are formed from the same cell type—endothelial cells. During embryonic development, individual endothelial cells grow and differentiate, into cells that will then become arteries, and those that will become veins.

Yet precisely what drives this process has long eluded researchers. Now, scientists at the Gladstone Institutes have identified the molecular signals that direct it, and in so doing, illustrate how even the most complex of biological systems can be directed by the most subtle shifts in molecular signaling.

In the latest issue of [*Developmental Cell*](#), researchers in the laboratory of cardiovascular researcher [Benoit Bruneau, PhD](#), describe the precise order and timing of signals that spur the formation of arteries. Specifically, they piece together a molecular signaling pathway by which a protein called *vascular endothelial growth factor* (Vegf) directs the activation of *Delta-like 4* (Dll4), which is critical to artery formation.

Vegf: the man behind the curtain

Arteries and veins each have different identities and distinct functions. Arteries carry oxygenated blood from the heart out to tissues, while veins carry unoxygenated blood back to the heart. Understanding how arteries are made at the molecular level—and specifically how they differ from veins—is important not only for understanding diseases in which arteries and veins connect abnormally, but also to inform strategies for making new arteries, which could prove invaluable for treating coronary artery disease.

The key to this understanding lies with Dll4, one of the earliest known genes involved in artery formation. In fact, scientists currently use Dll4 as a marker to identify which cells will grow and differentiate into arteries, and which will not. Dll4 works by binding to another protein—known as Notch—which in turn promotes artery formation.

But there is a third player in this process: Vegf. Vegf is a growth factor secreted at discreet time points and locations in the developing embryo to support proper blood vessel formation.

In addition to its receptor targets, it also activates Dll4. But exactly how Dll4, Notch and Vegf all work in concert to transform early embryonic cells into cells that form arteries has stumped researchers.

“We knew that Dll4, when activated, directs artery formation, but couldn’t pinpoint how it is activated in the first place,” said Dr. Bruneau, who is also a professor of pediatrics at the University of California, San Francisco, with which Gladstone is affiliated. “Here, we’ve mapped the series of steps that *precede* Dll4 activation and that set the stage for the formation of arteries—shedding light into a so-called ‘black box’ of embryonic development.”

In this study, the team delved deep into the nucleus of cells belonging to mouse and zebrafish embryos—two important animal models of embryonic development—in order to determine how the Dll4 gene is turned on. They used sophisticated molecular biology approaches, together with experiments that deleted specific genes from the animal models, to fill in the steps that led from Vegf signaling to Dll4 activation. And what they found was surprising.

“Vegf sets off a signaling cascade that eventually activates a group of proteins, called *Mitogen Activated Protein Kinases*, or MAPKs,” said Joshua Wythe, PhD, the paper’s lead author. “This, in turn, activates another group of proteins, called *ETS transcription factors*, and it is this signaling relay—from Vegf to MAPKs to ETS factors—that turns on and maintains Dll4 activity, helping the arterial cells grow and gain their cellular identity over time.”

In biology, a signaling cascade is a series of chain reactions that help cells amplify a particular signal—similar to a domino effect. Here, the team identified Vegf as being the first domino, followed by the activation of MAPKs, then ETS factors and so on.

“Interestingly, the ETS factors aren’t specific to soon-to-be arterial cells, but rather they are present throughout the embryo,” explained Jason Fish, PhD, from the University of Toronto, who collaborated with the Gladstone team. “Instead, the Vegf signaling cascade alerts *only* those MAPKs and ETS factors within the realm of Dll4—assuring only the correct cells grow and differentiate over time to form arteries.”

This research is important not only because it uncovers the molecular link between Vegf and Dll4, but also because it shows how signaling cascades like this one can direct genes—which are normally active throughout the embryo—to perform specific tasks only in specific cell types.

“In the future we will refine our approach to see whether this signaling cascade regulates other arterial genes in the developing embryo,” said Dr. Bruneau. “We hope this research will help inform clinicians into congenital defects related to the formation and maintenance of arteries and veins, and may also yield new strategies that can coax the development of arteries from stem cells—which may prove useful for treating coronary artery disease.”