

From Sci Fi to reality: Unlocking the secret to growing new limbs

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The axolotl, or Mexican salamander, is one of the three regenerative species described in a new paper that identified common genetic regulators governing limb regeneration in all three species. The findings suggest that these regulators aren't specific to individual species, but have been conserved through evolution, including in humans. The discovery opens the door to being able to manipulate these regulators with drug therapies. Credit: MDI Biological Laboratory

Many lower organisms retain the miraculous ability to regenerate form and function of almost any tissue after injury. Humans share many of our genes with these organisms, but our capacity for regeneration is limited. Scientists at the MDI Biological Laboratory in Bar Harbor, Maine, are studying the genetics of these organisms to find out how regenerative mechanisms might be activated in humans.



The ability of animals to regenerate body parts has fascinated scientists since the time of Aristotle. But until the advent of sophisticated tools for genetic and computational analysis, scientists had no way of studying the genetic machinery that enables <u>regeneration</u>. Using such tools, scientists at the MDI Biological Laboratory have identified genetic regulators governing regeneration that are common across species.

In a paper published in the journal *PLOS ONE*, MDI Biological Laboratory scientists Benjamin L. King, Ph.D., and Voot P. Yin, Ph.D., identified these common genetic regulators in three regenerative species: the zebrafish, a common aquarium fish originally from India; the axolotl, a salamander native to the lakes of Mexico; and the bichir, a ray-finned fish from Africa.

The discovery of genetic mechanisms common to all three of these species, which diverged on the evolutionary tree about 420 million years ago, suggests that these mechanisms aren't specific to individual species, but have been conserved by nature through evolution.

"I remember that day very well—it was a fantastic feeling," said King of the discovery. "We didn't expect the patterns of genetic expression to be vastly different in the three species, but it was amazing to see that they were consistently the same."

The discovery of the common genetic regulators is expected to serve as a platform to inform new hypotheses about the genetic mechanisms underlying <u>limb regeneration</u>. The discovery also represents a major advance in understanding why many tissues in humans, including limb tissue, regenerate poorly—and in being able to possibly manipulate those mechanisms with drug therapies.

"Limb regeneration in humans may sound like science fiction, but it's within the realm of possibility," said Yin. "The fact that we've identified



a genetic signature for limb regeneration in three different species with three different types of appendages suggests that nature has created a common genetic instruction manual governing regeneration that may be shared by all forms of animal life, including humans."

In particular, the scientists studied the formation of a mass of cells called a blastema that serves as a reservoir for regenerating tissues. The formation of a blastema is the critical first step in the <u>regeneration</u> <u>process</u>. Using sophisticated genetic sequencing technology, King and Yin identified a common set of genes that are controlled by a shared network of genetic regulators known as microRNAs.

"Scientists here are studying an evolutionarily diverse range of animals to gain insight into the genetic mechanisms underlying the repair and regeneration of complex tissues and why these processes are poorly active in humans," said Kevin Strange, Ph.D., the laboratory's president. "The value of our approach is confirmed by this remarkable study, which for the first time reveals a <u>genetic</u> network governing limb regeneration that is common across three evolutionarily distinct animal species."

The study also has implications for wound healing, which, because it also requires the replacement of lost or damaged tissues, involves similar genetic mechanisms. With a greater understanding of these mechanisms, treatments could potentially be developed to speed wound healing, thus reducing pain, decreasing risk of infection and getting patients back on their feet more quickly.

Another potential application is development of more sophisticated prosthetic devices. When a limb is amputated the nerves at the site of amputation can be damaged. The repair and regeneration of these nerves could potentially enable the development of more sophisticated prostheses that could interface with these nerves, allowing for greater



control.

While speedier wound-healing and improved prostheses may be on the nearer-term horizon, the ability to regrow limbs is a long way off. How long? "It depends on the pace of discovery, which is heavily dependent on funding," Yin said. He predicted the timeline could be hastened if enough research funding were available. "Unfortunately," he added, "we are in a period of greatly diminished funding for scientific research."

More information: Benjamin L. King et al. A Conserved MicroRNA Regulatory Circuit Is Differentially Controlled during Limb/Appendage Regeneration, *PLOS ONE* (2016). <u>DOI: 10.1371/journal.pone.0157106</u>

Provided by Mount Desert Island Biological Laboratory

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