

## From growing roots, clues to how stem cells decide their fate



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Long-term 4D confocal imaging of SHR reveals dynamics inconsistent with bistability. **a**, Diagram of *Arabidopsis* wild-type and *SHR:GAL4–GR UAS:SHR–GFP shr2* mutant roots showing proliferative and formative division planes (adapted from ref. <sup>52</sup>). SHR moves from the central tissues of the root into the adjacent cell layer. *SHR* expression and formative divisions occur in the inducible line upon treatment with dex. Yellow, QC (quiescent center); orange, CEI (cortex–endodermal initial); red, CEID (cortex–endodermal initial daughter) and *shr* mutant layer; blue, cortex; purple, endodermis. **b**, Diagram of



the SHR-SCR regulatory network controlling formative division based on Cruz–Ramirez et al.<sup>3</sup>. c, Confocal median longitudinal sections showing GFPlabeled SHR and H2B–RFP at timepoints after induction with 10 µM dex. Images are representative of independent timecourse experiments with eight roots. Numbers at the top left show the first five cell positions in the mutant ground tissue. Gamma is set to 0.75 to show signal in the mutant layer for the GFP-only images. Top and bottom show different roots. White arrows, formative divisions. Scale bars, 50 µm. d, Raw (gray) and smoothed (green) SHR trajectory (SHR-GFP/H2B-RFP fluorescence intensity) over time in the first five cells of a single cell file after full induction (10  $\mu$ M dex). Plots are representative of 211 cells from independent time courses with 8 roots. Possible low and high steady states are indicated for cell 1. Black dashed line, proliferative division; orange dashed line, formative division. a.u., arbitrary units. e, SHR trajectory predicted by the Cruz-Ramirez model showing low and high steady states. f, SHR trajectories for cells that show a low peak of SHR accumulation hours prior to dividing formatively. Roots were treated with low dex (0.02  $\mu$ M or 0.03  $\mu$ M). Dark green, SHR trajectory corresponding to images in g. g, Median longitudinal sections through a root tip treated with low dex (0.02 µM) highlighting a cell with a low transient peak of SHR prior to dividing formatively. Plots and images in **f** and **g** are representative of 15 cells from 10 roots showing similar behavior. Scale bars, 10 µm.

It might look like a comet or a shooting star, but this time-lapse video is actually a tiny plant root, not much thicker than a human hair, magnified hundreds of times as it grows under the microscope.

Researchers at Duke University have been making such movies by peering at <u>stem cells</u> near the root's tip and taking snapshots as they divide and multiply over time, using a technique called light sheet microscopy.

The work offers more than a front-row seat to the drama of growing roots. By watching how the cells divide in response to certain chemical



signals, the team is finding new clues to how stem cells choose one developmental path over another.

The research could also point to new ways to prevent stem cell division from going awry, as happens in cancer and other diseases.

The team's latest results appeared Jan. 31 in the journal Nature.

The work touches on a fundamental question in biology, said associate research professor Cara Winter: "How do cells acquire their identities?" In other words, "how do you get all of the various cell types that make up an organism?"

Just as the human body is made up of many different kinds of cells—in the brain, muscles, bones and elsewhere—plants, too, contain various cell types specialized for different tasks.

Whether in the roots, branches, flowers or leaves, virtually all tissues of a plant descend from small groups of unspecialized stem cells that produce new cells by dividing.

Each time a stem cell divides, it faces a choice: it can either produce two new stem cells like itself, or it can make one copy of itself plus one cell that will branch off to become something new.

It's the latter process, known as asymmetric division, that generates the myriad cell types needed to form a complex organism like a plant, or a human being.

An obvious question then is: How do dividing stem cells choose one path over the other?

This was the question driving Winter and co-first author Pablo Szekely,



both researchers in the lab of late biologist Philip Benfey of Duke, as they watched days of root growth in Arabidopsis thaliana, a spindly member of the mustard family.

The researchers focused on two key regulators of cell division in Arabidopsis—proteins called short-root and scarecrow that, together, prompt dividing <u>root</u> cells to make the switch.

By labeling these proteins with glowing fluorescent tags, they were able to track the activity of the proteins and their effects on dividing stem cells in real time. Light-sheet microscopy allowed them to peer inside the roots' translucent tissues for up to 50 hours without harming them.

Counter to previous predictions, the researchers showed that even low levels of these proteins, present early in the process of one cell becoming two, are enough to trigger a switch to asymmetric division.

"All they have to do is reach a certain threshold," said Szekely, who joined the Benfey lab as a postdoctoral researcher in 2020.

The findings have implications for humans and other animals too, the researchers said.

That's because, although plants and animals <u>diverged</u> more than a billion years ago, they inherited much of the same basic molecular tool kit—including many of the same "housekeeping" genes that are necessary for cells to function.

The same genes that regulate cell division in plants like Arabidopsis perform similar jobs in animals, including humans. <u>Previous research</u> shows that when asymmetric division is disrupted, cells can multiply out of control and form tumors.



"Cells need to have a program during development: first divide like this, then divide like that," Szekely said. "It has to be tightly regulated in order for everything to work."

**More information:** Cara M. Winter et al, SHR and SCR coordinate root patterning and growth early in the cell cycle, *Nature* (2024). DOI: 10.1038/s41586-023-06971-z

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