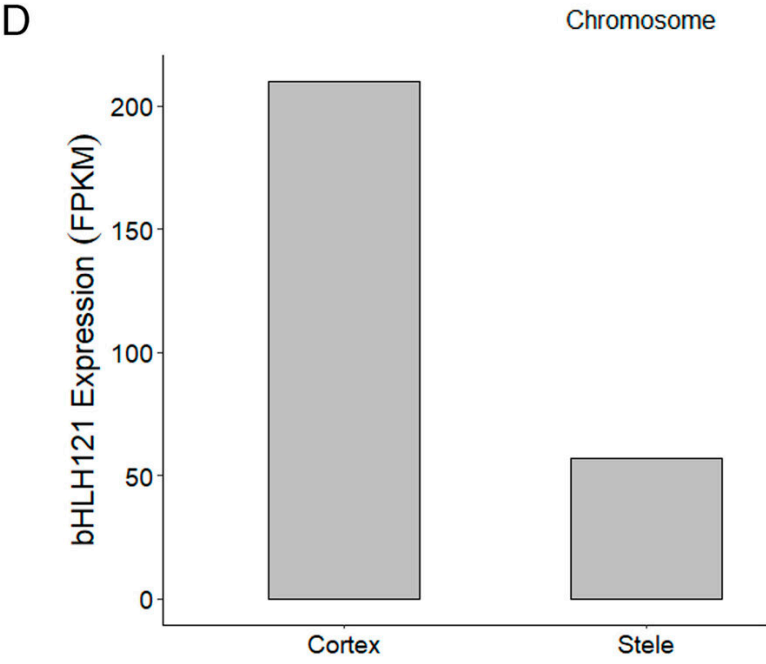
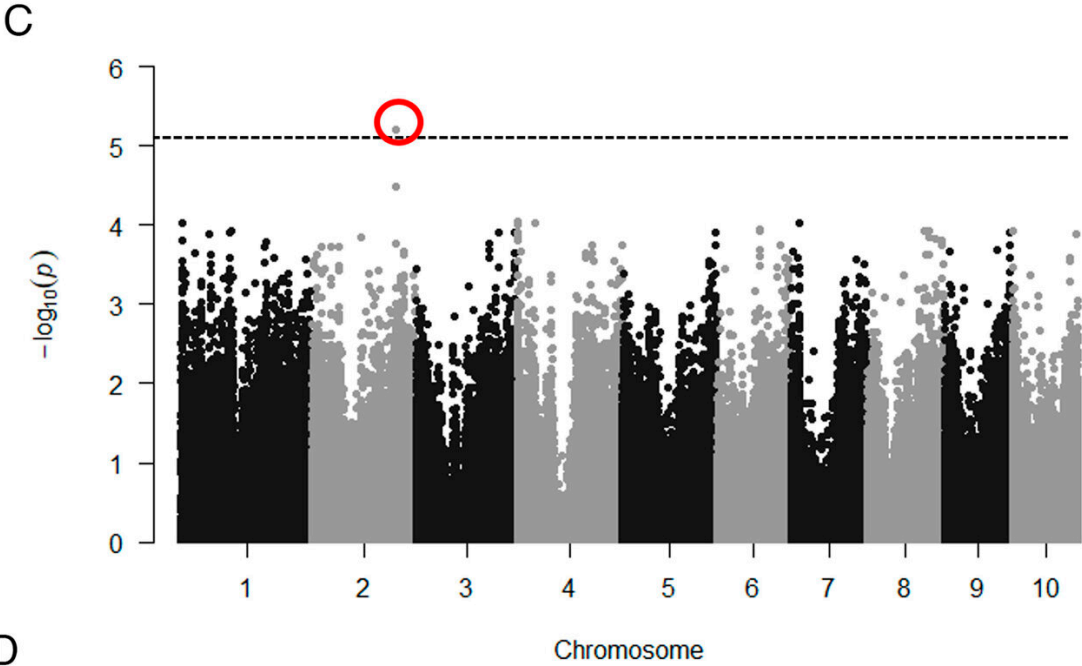
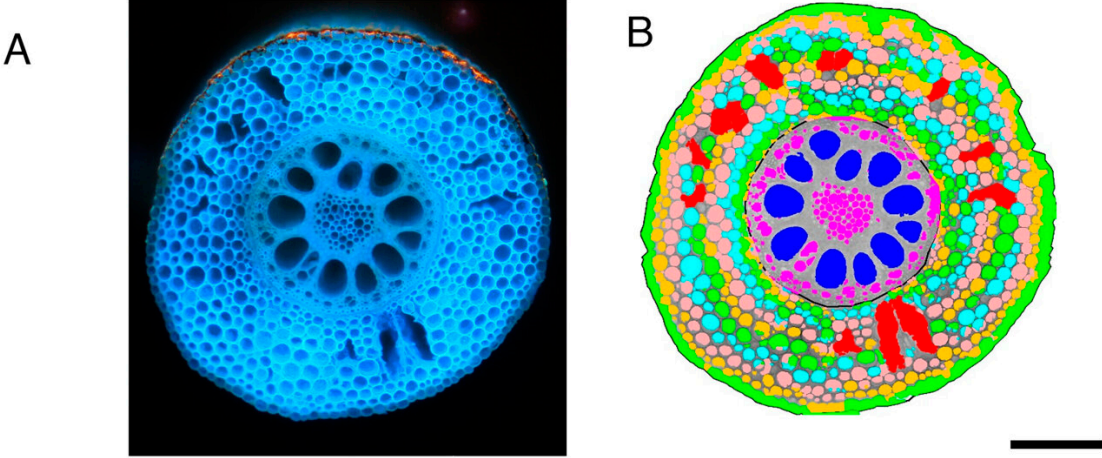


# Discovery of novel gene to aid breeding of climate resilient crops

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Identification of mutant locus associated with root cortical aerenchyma in maize roots. (A) An image of root cross-sectional surface captured by laser ablation tomography (LAT). (B) Root anatomical phenotypes were quantified by *RootScan* software. (C) Manhattan plot of GWAS results for root cortical aerenchyma. Chromosome-wide significance threshold (horizontal line) was set using the simpleM method ( $-\log_{10}p = 5.32$ ). Significant SNPs were located in gene model GRMZM2G083504 (*ZmbHLH121*) encoding a ZmbHLH121 transcription factor. (D) Relative expression of *ZmbHLH121* in the stele and cortex of a primary root from a 3-d-old maize seedling. (Scale bar, 100  $\mu\text{m}$ ).

Researchers have revealed for the first time how a key gene in plants allows them to use their energy more efficiently, enabling them to grow more roots and capture more water and nutrients.

An international team of plant scientists led by Penn State University and in collaboration with the University of Nottingham have discovered this novel regulatory gene (called bHLH121) that enables corn roots to acquire more water and nutrients. The findings have been published in the *Proceedings of the National Academy of Science*.

The gene controls the formation of air spaces among living root tissues (termed root cortical aerenchyma). Replacing a large percentage of root cells with airspaces saves the plant a lot of energy which is otherwise required to feed all these root cells. This makes roots metabolically more efficient, enabling them to use the resources saved to build more roots and explore the soil more effectively and capture more water and nutrients.

This discovery could lead to the breeding of crops that can withstand drought and low-nitrogen soil conditions and ultimately ease global food

insecurity, the researchers suggest.

"Identifying this gene and how it works will enable us to create more resilient crops that can withstand water and nutrient stress conditions being experienced as a result of [climate change](#)," says Rahul Bhosale, Assistant Professor in Crop Functional Genomics from the School of Biosciences and BBSRC Discovery Fellow.

The research team used powerful imaging tools developed in previous research at Penn State that rapidly measured cells in thousands of roots. An [imaging technique](#) called Laser Ablation Tomography was critical for this approach. This state-of-the-art approach is also now available at the University of Nottingham.

Hannah Schneider, Assistant Professor of Crop Physiology at Wageningen University & Research, Netherlands said, "We first performed the field experiments that went into this study starting in 2010, growing more than 500 lines of corn at sites in Pennsylvania, Arizona, Wisconsin and South Africa," she said. "I worked at all those locations. We saw convincing evidence that we had located a gene associated with root cortical aerenchyma."

This research revealed that mutant corn lines lacking the bHLH121 gene showed reduced root air space formation. In contrast, overexpressing bHLH121 caused more air space formation.

Characterization of these lines under suboptimal water and nitrogen availability in multiple locations revealed that the bHLH121 gene is required for [root](#) air space formation and provides a new tool for plant breeders to select varieties with improved soil exploration, and thus yield, under suboptimal conditions.

"These findings are the result of many people at Penn State and beyond

collaborating with us, working over many years," he said. "We discovered the function of the aerenchyma trait and then the gene associated with it, And, it came about because of technologies that have been devised here at Penn State, such as Shovelomics—digging up roots in the field—Laser Ablation Tomography and Anatomics Pipeline. We put all those together in this work," says Professor Jonathan Lynch, research lead at Penn State.

The results are significant, Lynch continued, because finding a gene behind an important trait that's going to help [plants](#) have better drought tolerance and better nitrogen and phosphorus capture looms large in the face of climate change.

"Those are super important qualities—both here in the U.S. and around the world," he said. "Droughts are the biggest risk to corn growers and are worsening with climate change, and nitrogen is the biggest cost of growing corn, from both a financial and environmental perspective. Breeding [corn](#) lines more efficient at scavenging for the nutrient would be a major development."

**More information:** Hannah M. Schneider et al, Transcription factor bHLH121 regulates root cortical aerenchyma formation in maize, *Proceedings of the National Academy of Sciences* (2023). [DOI: 10.1073/pnas.2219668120](#)

Provided by University of Nottingham

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