

# **Identifying traits early in development could predict future biomass production in *Miscanthus***

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Miscanthus sinensis. Credit: Lauren Quinn

New genetic tools have identified links between early establishment traits and biomass yield in *Miscanthus*. This marker-assisted selection technique could save years of waiting for *Miscanthus* plants to establish and become mature. The technique can be used to reliably select higher-yielding plants earlier in the *Miscanthus* life cycle.

The U.S. government has mandated a rise in domestic ethanol production from cellulosic bioenergy crops, spurring research to identify and improve high-yielding biomass crops. One of the strongest contenders is *Miscanthus*, a large Asian perennial grass. Several species and hybrids have proven to produce substantially greater biomass yields than switchgrass and other energy crops. However, *Miscanthus* crops can take from three to four years to establish before yields become reliable. A recent University of Illinois study used [genetic markers](#) to identify early developmental traits that correlate with future yield, potentially saving farmers and researchers years of time and effort.

According to the researchers, "this study begins to establish links between reproducible genetic markers and a number of key agronomic traits in *Miscanthus sinensis*."

The researchers measured developmental and biomass traits over a period of three establishment years in the offspring of a cross between *Miscanthus sinensis* cultivars 'Grosse Fontaine' and 'Undine.' They then extracted DNA from the plants and examined the resulting single nucleotide polymorphisms, or SNPs, to develop a genetic map. The technique improves upon older types of genetic markers that were not as tightly linked to particular genes controlling important biomass traits.



"It represents one of the very first maps that was made and it's also one of the first times we were able to map a number of genes associated with biomass productivity, and determine the locations of those genes in the *Miscanthus* genome," says U of I geneticist Jack Juvik.

On a practical level, the researchers saw strong positive correlations between biomass yield and plant basal circumference, height, and tiller (stem) number, suggesting that plants that are able to grow taller and produce more tillers in the first few years may achieve higher yields in the long term. Furthermore, there were negative correlations between flowering time and yield, with early flowering individuals producing less biomass. Breeders could make use of that information to improve early selection of plants with enhanced [biomass](#) productivity to accelerate the breeding program.

"The advantage to marker-assisted breeding is that you can grow seedlings, collect DNA, and probe for a large suite of DNA markers that are linked to genes that confer the characteristics you want. That can save a lot of time, because you can identify potential phenotypes without having to wait 3-4 years to get a mature plant," explained Juvik. "The value of this kind of system in *Miscanthus* is substantial in terms of breeding progress."

In addition to saving time and providing breeders with specific traits to look for in high-yielding plants, the techniques used in the study and the initial results establish a jumping-off point for future work.

Juvik notes, "This is the starting point. We've continued this work and applied it to other populations and to other questions about breeding *Miscanthus*. This sets up the foundation for moving into a range of different applications."

**More information:** Justin M. Gifford et al. Mapping the genome of

for QTL associated with biomass productivity , *GCB Bioenergy* (2015).  
[DOI: 10.1111/gcbb.12201](https://doi.org/10.1111/gcbb.12201)

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