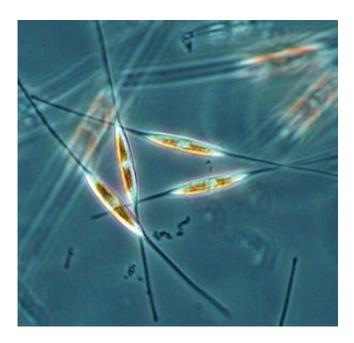


## In diatom, scientists find genes that may level engineering hurdle

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By manipulating the genes responsible for silica production in diatoms unicellular algae that encase themselves in intricately patterned, glass-like shells — scientists hope to produce faster computer chips. Photo: courtesy Wikimedia Commons

Denizens of oceans, lakes and even wet soil, diatoms are unicellular algae that encase themselves in intricately patterned, glass-like shells. Curiously, these tiny phytoplankton could be harboring the next big breakthrough in computer chips.



Diatoms build their hard cell walls by laying down submicron-sized lines of silica, a compound related to the key material of the semiconductor industry—silicon. "If we can genetically control that process, we would have a whole new way of performing the nanofabrication used to make computer chips," says Michael Sussman, a University of Wisconsin-Madison biochemistry professor and director of the UW-Madison's Biotechnology Center.

To that end, a team led by Sussman and diatom expert Virginia Armbrust of the University of Washington has reported finding a set of 75 genes specifically involved in silica bioprocessing in the diatom Thalassiosira pseudonana, as published today in the online Early Edition of the *Proceedings of the National Academy of Sciences*. Armbrust, an oceanography professor who studies the ecological role of diatoms, headed up the effort to sequence the genome of T. pseudonana, which was completed in 2004.

The new data will enable Sussman to start manipulating the genes responsible for silica production and potentially harness them to produce lines on computer chips. This could vastly increase chip speed, Sussman says, because diatoms are capable of producing lines much smaller than current technology allows.

"The semiconductor industry has been able to double the density of transistors on computer chips every few years. They've been doing that using photolithographic techniques for the past 30 years," explains Sussman. "But they are actually hitting a wall now because they're getting down to the resolution of visible light."

Before diatoms were appreciated for their engineering prowess, they interested ecologists for their role in the planet's carbon cycle. These photosynthetic cells soak up carbon dioxide and then fall to the ocean floor. They account for upwards of 20 percent of the carbon dioxide that



is removed from the atmosphere each year, an amount comparable to that removed by all of the planet's rainforests combined.

"We want to see which genes express under different environmental conditions because these organisms are so important in global carbon cycling," explains Thomas Mock, a postdoctoral researcher in Armbrust's lab and the paper's first author.

But research on these algae has uncovered other enticing possibilities. As he learned about diatoms, Sussman became intrigued by the fact that each species of diatom—there may be around 100,000 of them—is believed to sport a uniquely designed cell wall.

To determine which genes are involved in creating those distinctive patterns, the research team used a DNA chip developed by Sussman, UW-Madison electrical engineer Franco Cerrina and UW-Madison geneticist Fred Blattner, the three founders of the biotechnology company NimbleGen. Put simply, the chip allows scientists to see which genes are involved in a given cellular process. In this case, the chip identified genes that responded when diatoms were grown in low levels of silicic acid, the raw material they use to make silica.

Of the 30 genes that increased their expression the most during silicic acid starvation, 25 are completely new, displaying no similarities to known genes.

"Now we know which of the organism's 13,000 genes are most likely to be involved in silica bioprocessing. Now we can zero in on those top 30 genes and start genetically manipulating them and see what happens," says Sussman.

For his part, Sussman is optimistic that in the long run these findings will help him improve the DNA chip he helped develop — the very one used



to gather data for this research project. "It's like the Lion King song," he says. "You know, 'the circle of life."

Source: University of Wisconsin-Madison

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