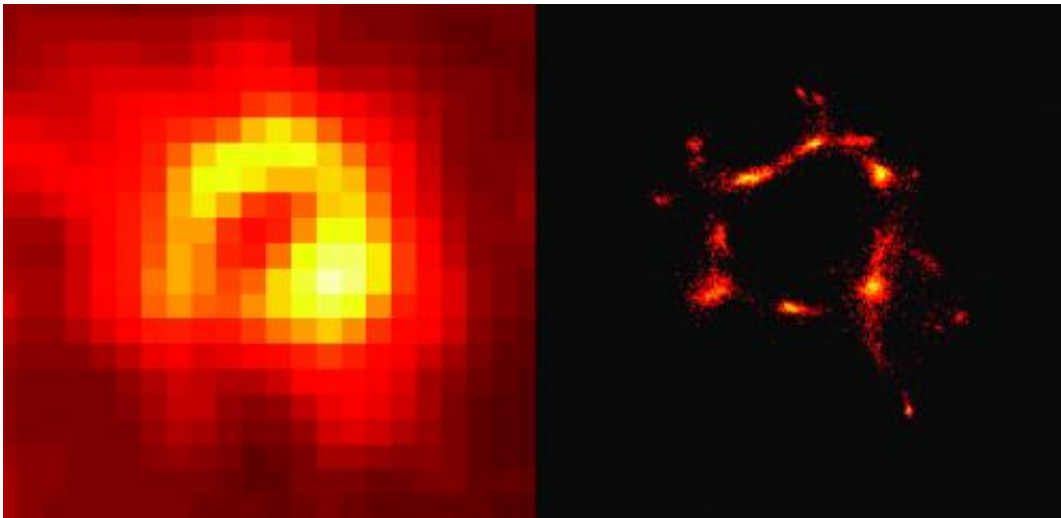


# Scientists use Amazon Cloud to view molecular machinery in remarkable detail

February 1 2013

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These microscope images of podosomes, cellular structures thought to be involved in cancer, show the differences in clarity produced by conventional microscopic techniques (left) and super-resolution imaging (right). Salk researchers have developed a method of utilizing cloud computing to significantly reduce the time necessary for processing super-resolution images. Credit: Salk Institute for Biological Studies

In this week's *Nature Methods*, Salk researchers share a how-to secret for biologists: code for Amazon Cloud that significantly reduces the time necessary to process data-intensive microscopic images.

The method promises to speed research into the underlying causes of

disease by making single-molecule microscopy of practical use for more laboratories.

"This is an extremely cost-effective way for labs to process super-resolution images," says Hu Cang, Salk assistant professor in the Waitt Advanced Biophotonics Center and coauthor of the paper. "Depending on the size of the data set, it can save over a week's worth of time."

The latest frontier in basic biomedical research is to better understand the "molecular machines" called proteins and enzymes. Determining how they interact is key to discovering cures for diseases. Simply put, finding new therapies is akin to troubleshooting a broken mechanical assembly line—if you know all the steps in the manufacturing process, it's much easier to identify the step where something went wrong. In the case of [human cells](#), some of the parts of the [assembly line](#) can be as small as single molecules.

Unfortunately, in the past conventional light microscopes could not clearly show objects as small as single molecules. The available alternatives, such as [electron microscopy](#), could not be effectively used with living cells.

In 1873, German physicist Ernst Abbe worked out the mathematics to improve resolution in light microscopes. But Abbe's calculations also established the optical version of the [sound barrier](#): the [diffraction limit](#), an unavoidable spreading of light. Think of how light fans out from a flashlight.

According to the Abbe limit, it is impossible to see the difference between any two objects if they are smaller than half the wavelength of the imaging light. Since the shortest wavelength we can see is around 400 nanometers (nm), that means anything 200 nm or below appears as a blurry spot. The challenge for biologists is that the molecules they want

to see are often only a few tens of nanometers in size.

"You have no idea how many single molecules are distributed within that blurry spot, so essential features and ideas remain obscure to you," says Jennifer Lippincott-Schwartz, a Salk non-resident fellow and coauthor on the paper.



This is Hu Cang, Assistant Professor, Waitt Advanced Biophotonics Center.  
Credit: Salk Institute for Biological Studies

In the early 2000s, several techniques were developed to break through the Abbe Limit, launching the new field of super-resolution microscopy. Among them was a method developed by Lippincott-Schwartz and her colleagues called Photoactivated Localization Microscopy, or PALM.

PALM, and its sister techniques, work because mathematics can see what the eye cannot: within the blurry spot, there are concentrations of photons that form bright peaks, which represent single molecules. The downside to these approaches is that it can take several hours to several days to crunch all the numbers required just to produce one usable image.

"It's like taking a movie, then you go through some very complex math, so what you see is the end result of processing, which is extremely slow because there's so many parameters," Cang says. "When I first saw PALM, I was shocked by how good it was. I wanted to use it right away, but when I actually tried to use it, I found its usefulness was limited by computing speed."

Even using statistical shortcuts, processing these images was still so intense that a supercomputer was required to reduce the time to a practical level. "Calculating an area of 50 pixels can take nearly a full day on a state-of-the-art desktop computer," says Lippincott-Schwartz. "But what you'll have achieved is the difference between a guess and a definitive answer."

In their [Nature Methods](#) paper, the researchers offer other scientists the tools they need to use an easier alternative—the Amazon Elastic Compute Cloud (Amazon Elastic EC2), a service that provides access to supercomputing via the Internet, allowing massive computing tasks to be distributed over banks of computers.

To make PALM more practical for use in biomedical research, the team

wrote a computer script that allows any biologist to upload and process PALM images using Amazon Cloud.

As a demonstration, Cang, Lippincott-Schwartz and post-doctoral researcher Ying Hu reconstructed the images of podosomes, which are molecular machines that appear to encourage cancer cells to spread. In one instance, they dropped the time needed to process an image from a whole day to 72 minutes. They also imaged tubulin, a protein essential for building various structures within cells. In that case, they were able to drop the time from nine days to under three and a half hours.

Their new paper provides a how-to tutorial for using the code to process PALM images through Amazon Cloud, helping the other labs achieve similar increases in speed.

Provided by Salk Institute

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