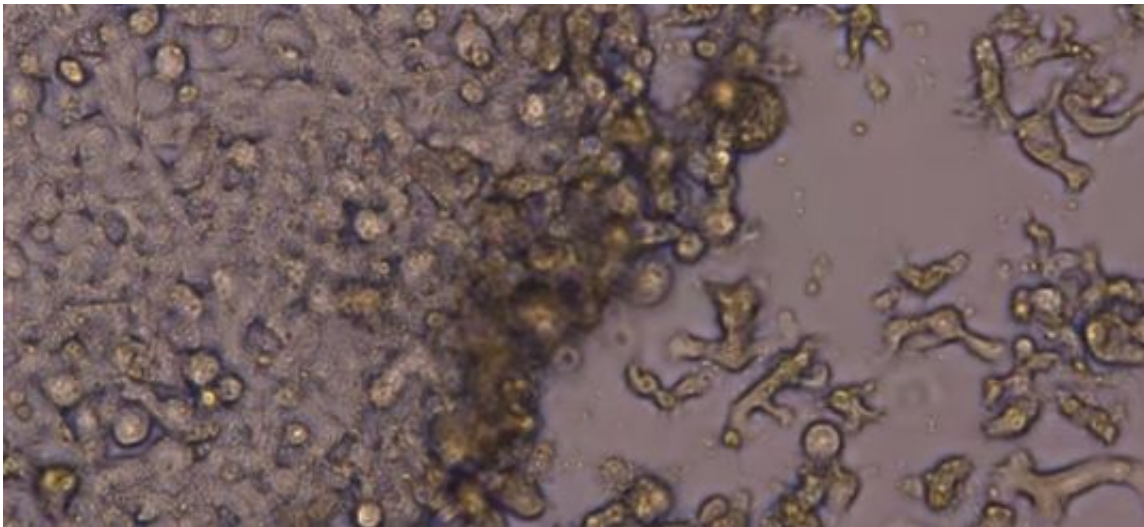


Next-generation genomic tests ID brain-eating amoeba

September 10 2015, by Nicholas Weiler



Last summer, a 74-year-old resident of San Francisco's Chinatown was admitted to San Francisco General Hospital and Trauma Center with fever and disorientation, which rapidly degenerated into a coma.

Her symptoms screamed that she was suffering from encephalitis, an infection of the brain, but a series of tests and drugs—for tuberculosis, bacterial and fungal infections, even toxoplasmosis—could neither identify the cause of infection nor do anything to resuscitate her.

She died 16 days later, her illness still a mystery to her clinicians.

Sadly, this outcome is all too common in cases of brain infection, which go unsolved six times out of 10 in California, according to a 2006 UC San Francisco study.

Biochemist Joe DeRisi, Ph.D., and neurologist Michael Wilson, M.D., of UCSF want to change that statistic, and they think they have the tools to do it.

When the researchers learned of last summer's case of mystery encephalitis, they turned to cutting-edge genomic tools to track down the cause of the woman's disease.

They using rapid next-generation genomic sequencing to search the patient's spinal fluid for non-human DNA, then compared what they found against databases of genes of every known invasive microorganism or virus. This "metagenomic" approach rapidly found a match: a common soil-dwelling [amoeba](#) called *Balamuthia mandrillaris*, as the researchers reported last month in the journal *Annals of Neurology*.

The researchers and their colleagues have recently set up a new Center for Next-Gen Precision Medicine Diagnostics to help hospitals make genome-based screens for the root causes of encephalitis part of their standard of care.

The technology could save money as well as lives. Over her two-week hospitalization, care for last summer's *Balamuthia* victim cost more than \$100,000. The metagenomic screen, by comparison costs a mere \$2,000—and typically takes two days.

One of the center's first priorities will be a study to determine when in the course of an encephalitis patient's care the technique would have the greatest clinical impact. Researchers will comb through databases of

encephalitis cases to identify when cheap, existing treatments like a \$100 blood draw are effective at diagnosing a patient's disease and when it's time for clinicians to turn to the center's big genomic guns.

"We all agree these metagenomic screens are happening too late," said UC Berkeley health economist Brent Fulton, PhD, MBA, part of the center's team who is leading the study. "When UCSF gets the call, doctors have already tried everything under the sun and it's a bad situation."

But how early should the test be done? It's still an open question, according to Fulton. "A \$2,000 test is nothing to sneeze at if it turns out to not change the usual course of care," he explained. "But balanced against a multi-week, half-million dollar hospitalization, it's nothing."

Through the new center, DeRisi hopes to make such screening a hospital's go-to test for mystery brain infections rather than a last resort. Currently doctors generally need to track down DeRisi and his peers directly to request help, he said. "We need to make this into an actual clinical diagnostic so a doctor who doesn't know my phone number can just order up this test."

Provided by University of California, San Francisco

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