

MinION Sketching, on the other hand, allows researchers to conduct cell line authentication in one afternoon. “We hope that the short time and DIY approach will reduce the burden and facilitate authentication,” Erlich said.

Neville Sanjana, a CRISPR biologist, is excited to make MinION Sketching a regular part of the work done in his lab. “There are very few techniques right now for cell-line authentication,” said Sanjana, a member at NYGC and assistant professor at NYU’s department of biology who was not involved in the research.

Sanjana said MinION Sketching is a step in the right direction for the field of clinical research. “This technique is one that is fast and more accurate than anything we have now. Cell authentication is really something that everyone should be doing.”

For the first time, the affordability of the MinION makes eventual widespread adoption of cell line authentication possible. The MinION is priced at \$1,000, with little additional costs to a laboratory already equipped to work with cancer cell lines. The Sketching code is available for free at github.com, and a link can be found in the online paper.

Creating a DNA Sketch

But, just how does MinION Sketching work? The entire workflow involved in reidentification of human samples involves collecting the sample, extracting the DNA, creating a library of strands to be sequenced, and sequencing using the MinION, which was created by Oxford Nanopore. The MinION is a nanopore sequencer containing a membrane embedded

with nanopore proteins that are just over a billionth of a millimeter wide. A steady ion current runs through the membrane. As single strands of DNA pass through the pores, each nucleotide (A, G, T, and C) is identified by the way its unique shape interrupts the ion flow.

The MinION generates readouts in real time of random pieces of DNA, while the Sketching software analyzes

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— **Neville Sanjana**, New York Genome Center and NYU Department of Biology

a selected number of single nucleotide polymorphisms, or SNPs, using a Bayesian algorithm. These variants make each person unique. “Every time a DNA read comes out, we are screening in semi-real time for variants that are informative and update the posterior probability. That gives us more strokes of the sketch, and we get better resolution of the match,” Zaaier said. Matches are made by comparisons to a database of known sequences. Researchers looking to use this method can download a SNP database, or, ideally, create their own.

In the paper, the researchers describe verifying their method by matching a strain of leukemic cells after three minutes of sequencing by the MinION and comparing it against a reference file in the Cancer Cell Line Encyclopedia database. The sketch required 91 SNPs to reach 99.9% accuracy. The researchers then contaminated that line with another and ran the test, again. The method correctly rejected a match if the contamination

level climbed above 25%.

This rapid procedure, along with the affordability of the MinION sequencer, promises to make cell-line authentication what it should be: standard procedure in every research lab. “No one wants to waste time and reagents working on the wrong cells,” Sanjana said. “At the right price, every lab will adopt this.” Zaaier added it also has the potential for a long list of

clinical applications. For example, it would take less than an hour before a surgery to verify the correct organ is being transplanted to a patient. Currently,

this verification takes a minimum of 24 hours. Likewise, it could be used in diagnostics in which patient samples are tested at various time points, making them more prone to mix-ups. “The speed of the MinION Sketching makes clinical sample authentication a game changer.”



Sophie Zaaier, Ph.D., of the Runway Startup Postdoc Program at the Jacobs Technion-Cornell Institute was the lead author of a study that could “democratize DNA fingerprinting.”