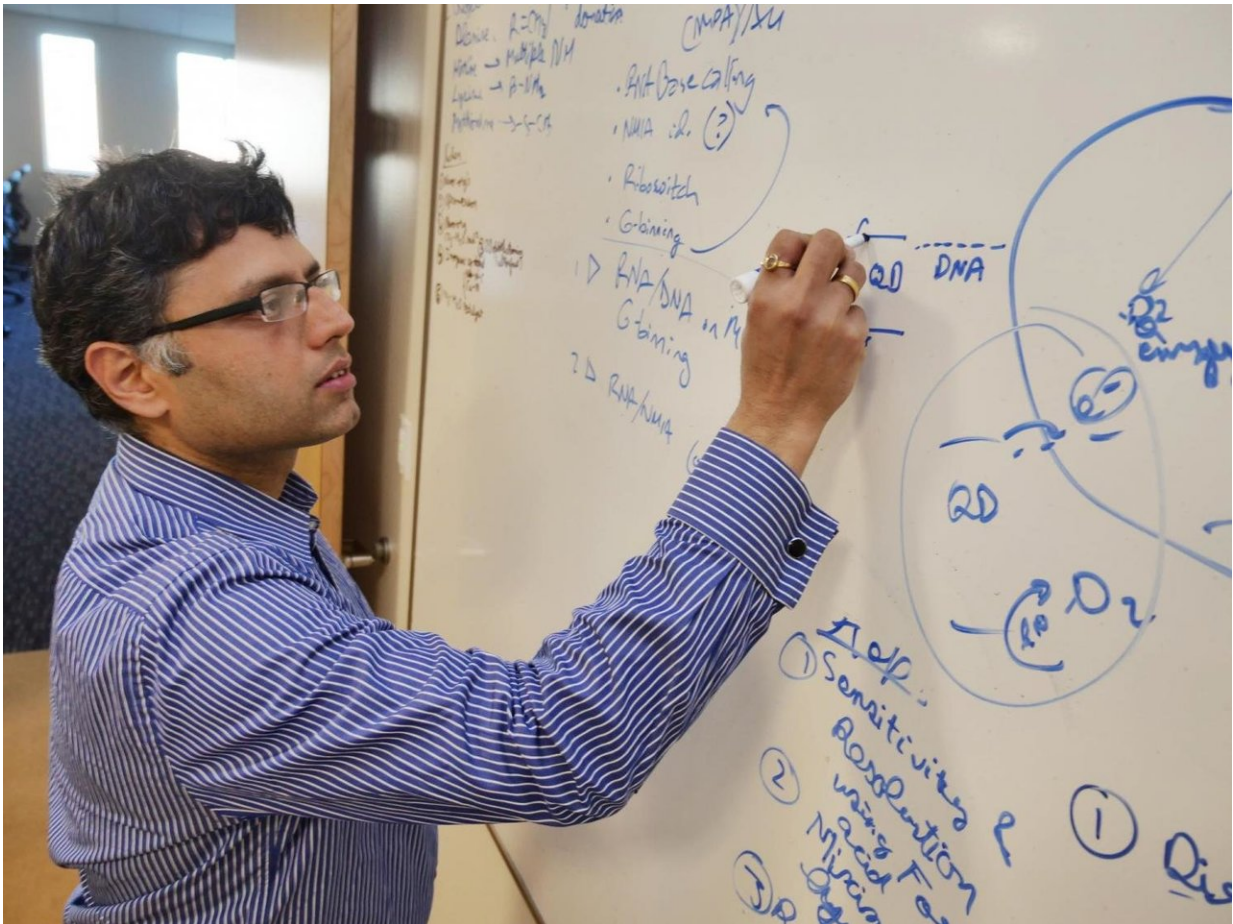


Rapid, cost-effective genetic screening within reach

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Assistant Professor Prashant Nagpal is researching new ways to create efficient, low-cost DNA sequencing that can provide important genetic information. Credit: Casey A. Cass / University of Colorado Boulder

CU Boulder researchers are developing new techniques for faster, more cost-effective single-molecule DNA sequencing that could have transformative impacts on genetic screening, paving the way for advances in vaccine development, early cancer detection and organ transplants.

The new methods—one of which is based in nanoscale quantum electronics and one in optics—could one day allow a person to sequence his or her genome in under two minutes using simple equipment, said Prashant Nagpal, an assistant professor in CU Boulder's Department of Chemical and Biological Engineering.

"We are exploring unique methods of providing accurate, useful genetic information faster and cheaper," said Nagpal.

DNA sequencing, which determines the precise order of nucleotides contained in biological molecules, can be used to map individual genes or entire genetic regions. Since its inception in the 1950s, sequencing has been incorporated into medicine, evolutionary biology and other bioscientific fields, all while increasing in speed and processing power by entire orders of magnitude.

Nagpal describes his new research as a timely reimagining of a technology that has existed for decades, but has run up against fundamental limitations in recent years.

"The human genome consists of over 3 billion DNA base pairs," said Nagpal. "Currently, sequencing relies on sampling 20 to 30 of these pairs at a time, which is good to have, but can only tell you so much."

The current sampling method, Nagpal says, opens the possibility for replication errors in the larger data set that might overlook fine-grain markers in a person's genetic profile. Single-molecule sampling, he says,

offers a more accurate picture.

Nagpal and his team turned to quantum electronics for the solution. They send a small charge through a DNA or RNA molecule, and record measurements of all nucleotides in single-molecules. The sequence identified the nucleotides with 99.7 percent accuracy, up from 80 percent achieved by other nanotechnology methods.

The advances in quantum-based sequencing were described recently in the journals *ACS Nano* and *Journal of the American Chemical Society*. The W.M. Keck Foundation provided support for all the single-molecule sequencing research.

Nagpal has also been developing an optics-based method for single-molecule sequencing that uses a diode laser and a camera comparable to that of an average smartphone to rapidly identify nucleotide sequences. Because of its unique physical properties as both a particle and a wave, light is a nimbler means of gathering information and transmitting it in compressed form.

In a forthcoming paper currently in press in the journal *Small*, Nagpal and his colleagues demonstrated a high-throughput sequencing method can be massively parallelized using a two-millimeter chip array that can take seconds instead of hours.

"With this block sequencing [method](#), you don't need to even sequence on the level of single letters like A, C, G and T," said Nagpal. "You can extrapolate the same information faster, similar to how you watch a compressed version of a movie on YouTube or Netflix."

The research is still in the early stages, and Nagpal says that the next step would be to apply the sequencing methods to more complex bacterial samples. In the future, he envisions a portable DIY sequencing kit that

would cost around \$1. A person could take a DNA swab, put it onto the microchip and wait 100 seconds to receive data that could reveal potentially risky genetic markers for cancer, for example, or provide crucial information for doctors attempting to match organs to transplant recipients.

"At scale, the potential applications are tremendous," said Nagpal. "From identifying antibiotic resistance to accelerating drug discovery, improvements in the speed and accuracy of sequencing could be transformative to many scientific fields."

Other co-authors on these single-molecule sequencing methods include Lee Korshoj, Sepideh Afsari, Manjunatha Sagar, Gary Abel and Assistant Professor Anushree Chatterjee, all of CU Boulder's Department of Chemical and Biological Engineering.

More information: Sepideh Afsari et al. Quantum Point Contact Single-Nucleotide Conductance for DNA and RNA Sequence Identification, *ACS Nano* (2017). [DOI: 10.1021/acsnano.7b05500](https://doi.org/10.1021/acsnano.7b05500)

Lee E. Korshoj et al. Conformational Smear Characterization and Binning of Single-Molecule Conductance Measurements for Enhanced Molecular Recognition, *Journal of the American Chemical Society* (2017). [DOI: 10.1021/jacs.7b08246](https://doi.org/10.1021/jacs.7b08246)

Provided by University of Colorado at Boulder

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