



Oxford Nanopore  
Technologies

## Revolutionising testing for critically ill patients

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Vorasuk Shotelersuk's goal is to enhance patient care through optimal genetic testing. Learn why Oxford Nanopore sequencing has the potential to provide the solution when precision and speed are paramount.

'To save the lives of critically ill patients, it is essential to have a rapid, high-yield, and practical diagnostic tool ... rapid [Oxford Nanopore] long-read genome sequencing is the solution available today'

**Professor Vorasuk Shotelersuk**  
Chulalongkorn University, Thailand



Reveal more biology to transform human health

# Ask bolder questions

Short-read sequencing is limited in its capacity for precise genetic diagnosis as detection of structural variants remains a challenge, and phasing information is inaccessible without parental samples<sup>2</sup>. Oxford Nanopore sequencing has the potential to address these two challenges<sup>2</sup>, whilst providing additional benefits in workload management and efficiency<sup>2,3</sup>.

## Ultra-rich data

61%

of cases for which  
causative variants  
were found<sup>2</sup>

9 days

median  
turnaround time<sup>2</sup>

8/18

cases where morbidity  
could have been prevented<sup>2</sup>

## Reveal more biology

Vorasuk's proof-of-concept study found that Oxford Nanopore sequencing has the potential to achieve a diagnostic yield of 61% (11/18), higher than the 46% (25/54) achieved with short-read exome sequencing in a quoted previous study<sup>2</sup>. Not only was nanopore sequencing more rapid, but the additional information meant that single nucleotide variants were reclassified, impacting treatment decisions<sup>2</sup>.

[This proof-of-concept study demonstrated the utility of singleton rapid \[Oxford Nanopore\] long-read genome sequencing as a first-tier diagnostic approach for critically ill patients with unknown causes<sup>2</sup>](#)



Read more about  
clinical research with  
Oxford Nanopore  
sequencing

### References

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[www.nanoporetech.com](http://www.nanoporetech.com)

phone +44 (0)845 034 7900

email [support@nanoporetech.com](mailto:support@nanoporetech.com)

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