

MinION GridION

Flexible, real-time, on-demand sequencing — in the lab or field

GridION



Ask bolder questions

Delivering any read length, ultra-rich datasets, and real-time insights, Oxford Nanopore sequencing answers the bigger, bolder research questions that you always wanted to ask. Welcome to sequencing without compromise.



Faster results

Richer insights

Near-sample, real-time workflows that don't require batching

captures more types of genetic variation

Highly accurate genomic data that

Accessible and affordable

Scalability that enables every application

What could you do with one MinION[™] Flow Cell?

Compatible with MinION and GridION[™] devices, MinION Flow Cells provide low-cost, scalable access to all the benefits of nanopore sequencing.

Any read length short to ultra long (20 bp to >4 Mb)

Direct methylation detection with native DNA or RNA sequencing

Multiplexed sequencing

of up to 96 samples with or without PCR



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SNPs &

phasing



Assembly

Structural variants

Make no compromises





Methylation



Full-length transcripts

Streamlined sample prep, on-demand sequencing, and real-time analysis for rapid results



1. Internal data generated using the Ultra-Long DNA Sequencing Kit.

Analyse

Bioinformatician



Command-line tools

- Access the latest algorithms
- Open-source tools developed by Oxford Nanopore or Nanopore Community
- Run pre-configured FPI2ME workflows

• Real-time results for time-critical applications

- Discover EPI2ME for streamlined, best practice analysis pipelines
- User-controlled run time stop sequencing when sufficient data generated, wash, and reuse flow cell
- Powerful GridION onboard compute alleviates the need for

MinION Mk1D

Your personal, portable DNA and RNA sequencer

A sequencer for everyone — empowering individual researchers, labs, and those new to DNA/RNA sequencing to perform in-house sequencing and take control of their timelines with a cost-effective, personal device. Weighing only 130 g and running off a laptop, MinION Mk1D generates tens of gigabases of real-time data in the field or lab.

Sample added to flow cell here

Sensor array with multiple nanopores for scaled-up sequencing

USB-C powered for fast data transfer; link to laptop or desktop computer to operate

Improved thermal tolerance for enhanced performance in the field



Indicator lights inform the user of run status and progress

Consumable flow cell where the biology and electronics come together for nanopore sequencing

Sensor chip works with custom ASIC for control and data acquisition

Compatible with MinION Flow Cells

GridION

Self-contained, easily deployable DNA/RNA benchtop nanopore sequencer

A flexible, self-contained, benchtop nanopore sequencer, running up to five MinION Flow Cells that can respond to the needs of multiple users on demand, across varied applications. Integrated, high-performance data processing alleviates the need for complex IT infrastructure.

Consumable flow cell where the biology and electronics come together for nanopore sequencing

Onboard data analysis offering real-time basecalling and adaptive sampling (on-device targeted sequencing)



GridION Q, part of the locked-down Q-Line range of devices for applied applications, also available. Find out more at nanoporetech.com/q-line.

Sample added to flow cell here

Five MinION Flow Cells can be operated individually or together, suitable for research labs and service providers

Expand your sequencing capabilities with PromethION™ 2 Solo run using GridION high-performance compute



Compatible with MinION Flow Cells

Choose your purchase plan





	MinION	GridION
Device	1	1
Flow cells	5	
Sequencing kits	1	
Wash kits	1	
Control kit	1	
Support*	1 year	1 year

* Extended support packages available — visit store.nanoporetech.com for more information.

Streamlined library preparation

A comprehensive range of library preparation kits are available, including direct, amplification-free DNA and RNA approaches that minimise potential bias and retain base modification information. Multiplexing (barcoding) options allow multiple samples to be analysed in a single sequencing run, maximising data generation while minimising costs.

	DNA		RNA		
Kit	Ligation	Rapid	PCR	Direct RNA	cDNA-PCR
Prep time	60 min	10 min	15 min + PCR	135 min	225 min + PCR
Input	~1 μg gDNA or 100–200 fmol amplicons	~200 ng gDNA or 50 ng for amplicons	1–5 ng gDNA	300 ng poly(A)+ RNA or 1 μg total RNA	10 ng poly(A)+ RNA or 500 ng total RNA
PCR required			\bigcirc		\bigcirc
Multiplexing	\bigcirc	\bigcirc	\bigcirc	In development	\bigcirc
Output	•••				• • •
Adaptive sampling	\bigcirc	\bigcirc	\bigcirc	In development	In development
Methylation included	\bigcirc	\bigcirc		\bigcirc	

Also available

- Ultra-Long DNA Sequencing Kit optimised for ultra-long DNA fragments and reads
- Application-specific library preparation kits (e.g. SARS-CoV-2 and 16S sequencing)
- Automatable workflows

View video tutorials, protocols, and more on the Nanopore Community: nanoporetech.com/support



Supporting your research at every step

In addition to the support included in your purchase plan and extensive online resources in the Nanopore Community, we offer personalised training courses to ensure successful optimisation of your nanopore sequencing projects.

	MinION Rapid Starter Training	GridION Advanced Training
Duration	2 days	2.5 days
Location	Online	On-site
Number of participants	Up to 2	Up to 4
Provided consumables	2 flow cells, 2 sequencing kits 6 flow cells, 2 sequencing kits	
User samples	1x control + 1x user sample 1x control + ≤4x user samples	
Content	The essentials of nanopore sequencing — from planning your experiment through to sequencing and an introduction to data analysis.	

Intuitive analysis with EPI2ME

The EPI2ME desktop application makes powerful genomic data analysis accessible to all scientists, regardless of bioinformatics expertise. Using an intuitive interface, users can navigate a growing range of open-source, best practice workflows that can be run on an Oxford Nanopore sequencing device*, laptop, desktop computer, cluster, or cloud service.

A rapidly growing range of workflows are available, including:

- Plasmid validation
- Adeno-associated virus verification
- Pathogen analysis: SARS-CoV-2, influenza, mpox, Mycobacterium tuberculosis
- Metagenomic species ID
- 16S-based microbial ID



* GridION or PromethION devices with integrated compute only.

- Antimicrobial resistance profiling
- Human variation: SVs, SNVs, and methylation including targeted approaches
- Transcriptomics: differential gene expression and transcript usage

Product specifications

MinION Mk1D device

- One flow cell position
- Up to 72-hour run time

Weight	Dimensions
130 g	W 55 mm H 13 mm D 125 mm

GridION device*

- Up to five individually addressable flow cells
- Up to 72-hour run time
- GPU-enabled real-time basecalling
- 7 TB SSD data storage
- 64 GB RAM
- Preloaded with Ubuntu OS and MinKNOW[™]

Weight Dimensions

W 370 mm | H 220 mm | D 365 mm 14.4 ka



Used for MinION and GridION devices

- Up to 72-hour run time
- Typical data output: 15-35 Gb
- Suitable applications include low-pass large genomes (e.g. human), whole prokaryotic genomes, metagenomics, targeted sequencing, large transcriptomes (cDNA), and small transcriptomes (direct RNA)
- Choose between DNA and direct RNA flow cells

Nanopore sequencing technology is advancing at an unprecedented pace, promising a future where portable sequencing will be routine in surveillance and many other fields.

> Jana Batovska La Trobe University

[With the GridION] we can have more than one flow cell starting at a different time, running different samples, running the same sample and don't forget you can multiplex on them as well.

> Dr Kim Judge Wellcome Sanger Institute

* Standard computer monitor, keyboard, and mouse required.



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