

Flexible, real-time, on-demand sequencing for your lab





[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.

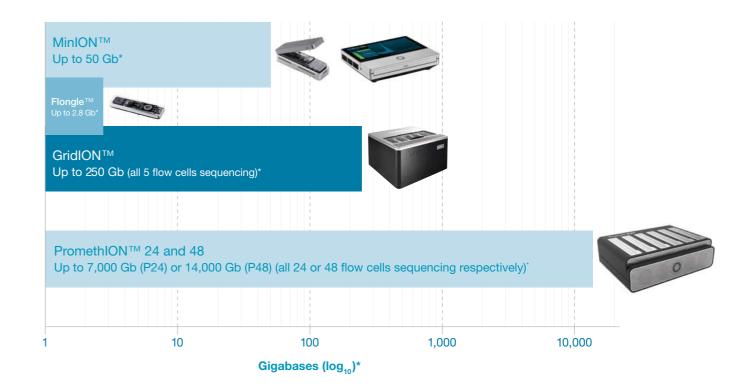
Dr. Kim Judge, Wellcome Sanger Institute

The GridION Mk1 from @nanopore is a breeze to install. It's about as plug and play as you can get with a piece of technology this advanced.

James Ferguson, Garvan Institute of Medical Research

GridION is a cost-effective and compact benchtop system offering on-demand sequencing with integrated real-time data processing

With the capacity to run five flow cells either concurrently or individually, GridION provides busy labs and service providers with cost-efficient access to the advantages of long-read, real-time nanopore sequencing. Integrated, high-performance data processing alleviates the need for complex IT infrastructure.



* Theoretical max output when system is run for 72 hours (or 16 hours for Flongle) at 420 bases / second Outputs may vary according to library type, run conditions, etc.

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



Prepare

- Streamlined library preps in as little as 10 minutes, with multiplexing options
- Scale according to your needs same chemistry and kits used for Flongle, MinION, GridION, and PromethION
- Run smaller sequencing tests and experiments or cost-effectively check your sample quality using Flongle on GridION

Sequence

- Sequence what you need, when you need it - no sample batching required
- Control each individual flow cell independently - run as many or as few as you wish at the same time, or add more whilst others are running
- GridION sequences DNA and RNA directly - meaning no amplification bias and retained modification (e.g. methylation) information

Analyse

- High-performance data processing capability with integrated basecall accelerator delivers real-time local analysis with no burden on existing IT infrastructure
- Discover EPI2ME and EPI2ME Labs for streamlined, best practice analysis pipelines and tutorials
- Choose to output the raw signal, or basecalled .fastq files, so you can use your own custom analysis pipelines

А	pplications include:	
	More information nanoporetech.com/products	
•	All the data, raw or basecalled, can be used in custom analysis pipelines written by the user for specific applications.	
•	Run open-source tools written and developed by the Nanopore Community. community.nanoporetech.com	
•	Explore your data and develop your bioinformatics skills with interactive, best practice workflows and tutorials. nanoporetech.com/analyse	
•	Use cloud-based or local EPI2ME platform for real-time analysis workflows. nanoporetech.com/analyse	
matic needed:		

- Flexible sequencing of whole genomes, targeted regions, and full-length RNA transcripts - all on one device
- Long reads enhance analysis of repetitive regions, structural variation, phasing, metagenomics, and more
- Quantify and characterise RNA splice variants, isoforms, and fusion transcripts

More information

nanoporetech.com/applications

Choose your GridION Mk1 plan



er Pack		CapEx*	
1		1	
60		-	
10		-	
5		-	
nonths		12 months	
luded		Included	
9,955		\$69,955	
	Buy nov	v store.nanoporetech.com	

Supporting your research at every step

Training options to suit your laboratory's needs

All GridION purchase plans include Remote Installation Assurance and expert support as standard — enabling easy device setup and complete optimisation of all your nanopore sequencing projects. Our technical specialists are available to answer all your questions on nanopore technology.

Remote Installation Assurance

- · Remote walk-through and an overview of nanopore sequencing with Q&A
- Remote installation and configuration assistance
- Remote platform QC check

Expert support

- Access to Nanopore Community support features
- End-to-end online protocol builder
- Customer Services initial setup call
- Technical Support experimental setup call
- Live online chat with Technical Support
- Facility to book Technical Support sessions as required

	Rapid Start Day Training	Advanced Nanopore Training
Location*	Oxford Nanopore labs, your site, or online	Oxford Nanopore labs, your site, or online
Duration	1 day	2 days
Participants	Private session with up to 2 participants	Private session with up to 4 participants
Content	Bespoke to suit your needs	Bespoke to suit your needs
User-provided samples processed	1 sample	Multiple samples
Flow cells included	2	6
Kits included	1–2†	2
Data analysis	Basic	Advanced
 * Third party reagents are provided only when training at Oxford Nanopore labs. * Depending on application. 	\$6,000	\$15,000

Nanopore Workshops

Choose the Nanopore Introduction Workshop and/or Data Analysis Overview to learn the essentials of nanopore sequencing and data analysis - from \$750 per person.

Simple plug-and-play setup

Product specifications







GridION Mk1*

- Up to 5 individually addressable flow cells
- 1 min 72 hour run time
- As much as 250 Gb⁺ total yield across the device
- 2,560 channels across the device can be sequencing at once
- GPU-based compute enabling real-time basecalling alongside additional analysis provided by EPI2ME, EPI2ME Labs, or a wide range of community-developed tools

Power requirements: 650 W

Storage: 4 TB SSD

Memory: 64 GB RAM

Weight: 11 kg

Dimensions: W 370 mm, H 220 mm, D 365 mm

Image courtesy of Dr. Alex Cagan Wellcome Sanger Institute

GridION Flow Cells

Same flow cells as MinION device.

- 1 min 72 hour run time
- Up to 50 Gb⁺ per flow cell
- 512 channels

Flongle

MinION/GridION flow cell adapter for more cost-effective sequencing of smaller tests and experiments.

- Up to 24 hour run time
- Up to 2.8 Gb⁺
- 126 channels

^{*} Standard computer monitor, keyboard, and mouse required.

[†] Theoretical max output when system is run for hours (or 16 hours for Flongle) at 420 bases / second. Outputs may vary according to library type, run conditions, etc.

