

Simply Fast

# FASTASeq

Speed redefined



#### Simply Fast

## **Swift**

2.0Hrs SE50

3.0Hrs SE100

6.8Hrs PE150

Lightning Chemistry Plus reduce cycle time

**50**%



# Saving

Reagent saving
Sequencing on demand

**Reduce reagent consumption** 

**70**%

# Simple

**User-friendly Interface** 

minutes
NGS run preparation



# Social Responsible

Smart waste sorting and processing design

RT shipping and stroage for load-and-go Reagent

#### **Product Parameter**

Parameter								
Flow cell type <sup>1</sup>	Lane	Throughput (Reads/FC)	Specification <sup>1</sup>	Read Length	Data Output²	Q30 <sup>2</sup>	TAT <sup>3</sup>	
FCL	1	20 M	SE100-D	SE100	2 Gb	≥90%	~ 3.0 hrs	
			100cycles	PE50			~ 4.0 hrs	
			300cycles	PE150	6 Gb	≥90%	~ 6.8 hrs	
	1	40 M	SE100-D	SE100	4 Gb	≥90%	~ 3.5 hrs	
FCM			100cycles	PE50			~ 4.5 hrs	
			300cycles	PE150	12 Gb	≥90%	~ 7.5 hrs	

Main Features				
Real time FastQ generation	Allow to resume from breakpoint			
Load-and-go reagent cartridge	RT shipping and storage for reagents			

<sup>1.</sup> FCL/FCM reagents available in Q3,2025, FCX reagents available in Q1 2026, FCH reagents available in Q2, 2026

#### **Application**

	Number of Samples / FC¹			
Application	FCL	FCM		
	20 M	40 M		
mNGS (based on probe capture) <sup>2 M Reads/sample</sup>	8	16		
mNGS (based on multiplex PCR) 0.5 M Reads/sample	32	64		
Small Oncology Panel (Tissue) 0.2 Gb/sample	24	48		
Small Oncology Panel (Plasma) <sup>2 Gb/sample</sup>	2	4		
HLA Typing <sup>0.2 Gb/sample</sup>	24	48		
PGT-A 5 M Reads/sample	3	6		
Microbial WGS <sup>1 Gb/sample</sup>	4	8		
NGS Library QC 0.1 M Reads/sample	160	320		

<sup>1.</sup> The above sample numbers for different application are only for reference. Users need to adjust the sample number according to the actual experiment.

<sup>2.</sup> This parameter is obtained based on the average of multiple test results of GeneMind P3 standard library. The date output and the proportion

of high quality data are affected by factors such as sample type, sample quality and effective flowcell utilization. The actual performance may vary.

3. The sequencing time to complete the sample read length plus the paired-end index (8+8), which included the time from sample loading to base calling and generate basefile.

## **Ordering Information**

PN	Sequencer
SQ00075	FASTASeq S Sequencing System Set
PN	Reagents
S000515	FASTASeq S Sequencing Kit V1.0 (FCL-D SE100-D)
S000516	FASTASeq S Sequencing Kit V1.0 (FCL 100cycles)
S000517	FASTASeq S Sequencing Kit V1.0 (FCL 300cycles)
S000518	FASTASeq S Sequencing Kit V1.0 (FCM-D SE100-D)
S000519	FASTASeq S Sequencing Kit V1.0 (FCM 100cycles)
S000520	FASTASeq S Sequencing Kit V1.0 (FCM 300cycles)

## **System Specification**

Dimensions (W $\times$ D $\times$ H)	626 mm×580 mm × 575 mm		
Net Weight	about 100 kg		
Power	≤1000 VA		
Power Requirements	100-240 V∼,50/60 Hz		
Operating Environment	Temperature:19°C-25°C Humidity:20%-80% (non-condensing) Altitude: below 3000 meter		
Computer Configuration	CPU:13th Gen Inter(R) Core(TM) i7-13700T Memory:64 GB Hard Drive:4 TB Operating system:Microsoft Windows 10		

## GeneMind Biosciences Co., Ltd.

Technical Support: **+86-400-822-3660** 

Website: en.genemind.com Email: info@genemind.com

Address: Room 502A/502B/602, Luohu Investment Holding Building, 112 Qingshuihe 1st Road,

Luohu District, Shenzhen, Guangdong, China





