

### You're driven by innovation. So are we.

The power of next-generation sequencing (NGS) has never been so expansive, promising, and exciting. Your boldest goals have every chance of being realized. At Illumina, we provide you with the tools and innovations you need to unlock the power of the genome.

In genetic disease, reproductive health, oncology, microbiology, agriculture, and beyond, researchers and clinicians are relying on Illumina systems to deliver data that powers groundbreaking insights.

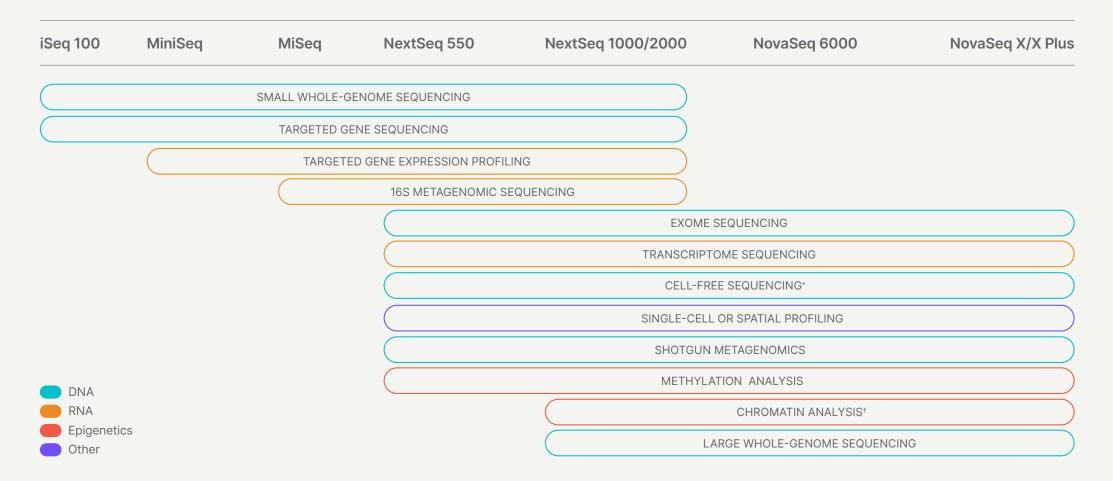
With a full suite of systems, we have the right solution to meet your ever-evolving needs.

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### A breadth of solutions. A whole world of answers.



<sup>\*</sup> Cell-free sequencing includes noninvasive prenatal testing (NIPT) and liquid biopsy.

<sup>†</sup> Chromatin analysis includes assay for transposase accessible chromatin (ATAC-Seq), chromatin immunoprecipitation (ChIP-Seq), and chromatin conformation capture (Hi-C).

From everyday tasks to your boldest projects, there's an Illumina system to meet your sequencing needs.‡

#### Research

Our benchtop sequencing solutions, from the  $iSeq^{TM}$  100 Sequencing System to the NextSeq<sup>TM</sup> 2000 Sequencing System, give you the power of Illumina NGS technology in a highly accessible and flexible design.

Our production-scale sequencing systems enable high-throughput, data-intensive applications. The NovaSeq $^{\text{TM}}$  X Series is quite simply a revolution in genomics, powering your studies with exceptional throughput and accuracy. Projects previously thought out of reach are now possible.

#### **Diagnostic**

For *in vitro* diagnostic (IVD) applications, clinical testing on the MiSeq™Dx,§ NextSeq 550Dx,§ and NovaSeq 6000Dx§ instruments leads to deep insights that help improve patient outcomes.



<sup>‡</sup> Throughput and data intensity determines system recommendations for methods and applications.

<sup>§</sup> For In Vitro Diagnostic Use. Not available in all regions and countries.

## NGS at your fingertips







| iSeq 100 System | MiniSeq <sup>™</sup>  | Syster |
|-----------------|-----------------------|--------|
|                 | A At all and a second |        |

| MiSeq Sy | ystem |
|----------|-------|
|----------|-------|

| Flow cell                | _                 | Mid-output        | Rapid      | High-output | Nano              | Micro      | v2            | v3         |
|--------------------------|-------------------|-------------------|------------|-------------|-------------------|------------|---------------|------------|
| Output range             | 144 Mb-1.2 Gb     | 2.1–2.4 Gb        | 2 Gb       | 1.65-7.5 Gb | 300–500 Mb        | 1.2 Gb     | 750 Mb-8.5 Gb | 3.8-15 Gb  |
| Single-end reads per run | 4M                | 8M                | 20M        | 25M         | 1M                | 4M         | 15M           | 25M        |
| Run time                 | 9–19 hr           | 17 hr             | < 5 hr     | 7–24 hr     | 17–28 hr          | 19 hr      | 5.5–39 hr     | 21–56 hr   |
| Maximum read length      | 2 × 150 bp        | 2 × 150 bp        | 1 × 100 bp | 2 × 150 bp  | 2 × 250 bp        | 2 × 150 bp | 2 × 250 bp    | 2 × 300 bp |
| Included data analysis   | Local Run Manager | Local Run Manager |            |             | Local Run Manager |            |               |            |



### Power and flexibility on your benchtop





#### NextSeq 550 System<sup>a</sup>

#### NextSeq 1000 and NextSeq 2000 Systems

| Flow cell                | Mid-output        | High-output | P1 <sup>b</sup>                   | P2 <sup>b</sup> | P3°        | P4 <sup>c</sup> |
|--------------------------|-------------------|-------------|-----------------------------------|-----------------|------------|-----------------|
| Output range             | 16-39 Gb          | 25-120 Gb   | 10-60 Gb                          | 40-240 Gb       | 120-360 Gb | 80-540 Gb       |
| Single-end reads per run | 130M              | 400M        | 100M                              | 400M            | 1.2B       | 1.8B            |
| Run time                 | 15–26 hr          | 11–29 hr    | 8–34 hr                           | 12-42 hr        | 18-40 hr   | 12-44 hr        |
| Maximum read length      | 2 × 150 bp        | 2 × 150 bp  | 2 × 300 bp                        | 2 × 300 bp      | 2 × 150 bp | 2 × 150 bp      |
| Included data analysis   | Local Run Manager |             | Onboard DRAGEN secondary analysis |                 |            |                 |

a. The NextSeq 550 System includes array scanning functionality for cytogenomic, methylation, and karyomapping applications.

b. Specifications for NextSeq 1000/2000 XLEAP-SBS™ reagents shown.

c. Specifications for NextSeq 2000 XLEAP-SBS reagents shown. P3 and P4 reagents are available for the NextSeq 2000 System only.



### Production-scale systems to maximize output







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| NovaSeq 6000 System            |               |                 |                   | NovaSeq X System |                |              | NovaSeq X Plus System |                 |            |            |
|--------------------------------|---------------|-----------------|-------------------|------------------|----------------|--------------|-----------------------|-----------------|------------|------------|
| Flow cell                      | SP            | S1              | S2                | S4               | 1.5B           | 10B          | 25B                   | 1.5B            | 10B        | 25B        |
| Flow cells processed per run   | 1 or 2        | 1 or 2          | 1 or 2            | 1 or 2           | 1              | 1            | 1                     | 1 or 2          | 1 or 2     | 1 or 2     |
| Output range                   | 65-<br>800 Gb | 134 Gb-<br>1 Tb | 333 Gb-<br>2.5 Tb | 280 Gb-<br>6 Tb  | 165–<br>500 Gb | 1–3 Tb       | 8 Tb                  | 165 Gb-<br>1 Tb | 1–6 Tb     | 8–16 Tb    |
| Single-end reads per flow cell | 800M          | 1.6B            | 4.1B              | 10B              | 1.6B           | 10B          | 26B                   | 1.6B            | 10B        | 26B        |
| Run time                       | 13–38 hr      | 13–25 hr        | 16–36 hr          | < 44 hr          | 17–23 hr       | 18–25 hr     | ~48 hr                | 17–23 hr        | 18–25 hr   | ~48 hr     |
| Maximum read length            | 2 × 250 bp    | 2 × 150 bp      | 2 × 150 bp        | 2 × 150 bp       | 2 × 150 bp     | 2 × 150 bp   | 2 × 150 bp            | 2 × 150 bp      | 2 × 150 bp | 2 × 150 bp |
| Included data analysis         | _             |                 |                   |                  | Onboard DR     | AGEN seconda | nry analysis          |                 |            |            |



### More clinical options. More meaningful answers.







| MiSeqDx Ins | strument |
|-------------|----------|
|-------------|----------|

NextSeq 550Dx Instrumenta

NovaSeq 6000Dx Instrumenta

|                                | •                          |  | •   |  |  | •   |   |  |
|--------------------------------|----------------------------|--|---|--|--|---|---|--|
|                                | MiSeqDx v3<br>(300 cycles) | Capabilities<br>in Research<br>mode <sup>a</sup> | NextSeq 550Dx<br>High Output v2.5<br>(300 cycles) | NextSeq 550Dx<br>High Output v2.5<br>(75 cycles) | Capabilities<br>in Research<br>mode <sup>a</sup> | NovaSeq 6000Dx<br>S2 v1.5<br>(300 cycles) | NovaSeq 6000Dx<br>S4 v1.5<br>(300 cycles) | Capabilities<br>in Research<br>mode <sup>a</sup> |
| Flow cells processed per run   | 1                          | 1  | 1   | 1  | 1  | 1 or 2                                    | 1 or 2                                    | 1 or 2   |
| Output range                   | ≥ 5 Gb                     | 300 Mb-15 Gb                                     | ≥ 90 Gb   | ≥ 22.5 Gb  | 16-120 Gb  | 1–2 Tb                                    | 3–6 Tb                                    | 80 Gb-6 Tb                                       |
| Single-end reads per flow cell | ≥ 15M                      | 25M  | ≥ 300M  | 400M   | 400M   | 4.1B                                      | 10B                                       | 10B  |
| Run time                       | 24 hr                      | 5.5–56 hr  | < 35 hr   | < 11 hr  | 11–29 hr   | ≤ 40 hr                                   | ≤ 45 hr                                   | 13–44 hr   |
| Maximum read length            | 2 × 150 bp <sup>c</sup>    | 2 × 300 bpb                                      | 2 × 150 bp  | 1 × 75 bp  | 2 × 150 bp                                       | 2 × 150 bp                                | 2 × 150 bp                                | 2 × 250 bp                                       |
| Included data analysis         | Local Run Ma               | nager  | Local Run Manager                                 |  |  | Paired DRAGEN se                          | rver                                      |  |

a. In Research (RUO) mode, MiSeqDx, NextSeq 550Dx, and NovaSeq 6000Dx instruments have the same performance specifications as the MiSeq, NextSeq 550, and NovaSeq 6000 Systems, respectively.

b. Refer to package insert for assay-dependent specifications.





# Connecting data to insights

With an Illumina system, efficiencies are built in. Our comprehensive software solutions help reduce bioinformatics bottlenecks and streamline your genomics workflow. Whether you're just getting started, or you're in rapid scale mode, Illumina Connected Software\*\* unlocks the power of your data in applications spanning oncology, rare disease, and infectious disease.

Integrated with our sequencing systems, Illumina Connected Software supports genomic and clinical researchers from primary to tertiary analysis, optimizes lab and sample management, and accurately calls genetic variations. Balancing approachability with customization, Illumina Connected Software enables insights for single-sample or population-wide studies.

Meeting you where your data are, Illumina offers solutions for both local and cloud analysis. We are committed to relentless innovation, creating new bioinformatics technologies that expand access to genomics for all.

#### **Proven accuracy**

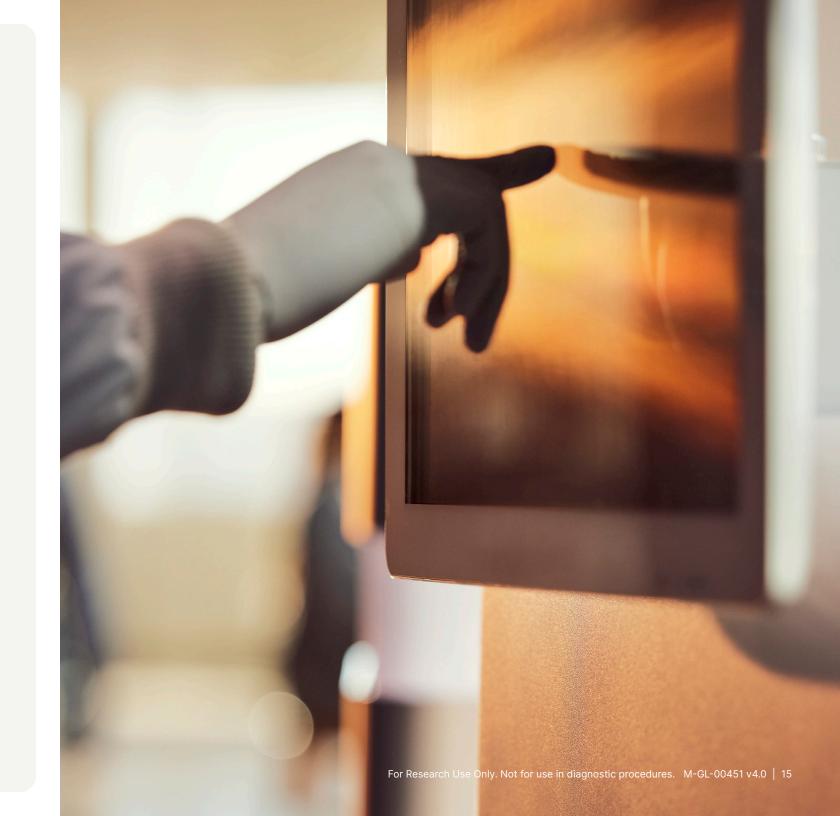
Highly accurate sequencing by synthesis (SBS) chemistry plus DRAGEN™ secondary analysis deliver award-winning germline and somatic variant calling.†† With onboard DRAGEN analysis available on select instruments, users can gain significant cost savings for accurate, comprehensive, and efficient NGS analysis.

#### **High standards for data privacy**

To meet the most stringent security requirements, our software products are built with security and compliance at the core. Data sharing security and governance, audit trails with encryption, and controlled sharing ensure your data are kept safe and secure.

#### **Trusted technology partners**

Dedicated to your success, the Illumina Informatics Services team brings bioinformaticians, data scientists, and designers to help you customize and optimize your analysis workflow and minimize your development burden.





### Support that never stops

For Illumina, innovation doesn't stop at developing best-in-class systems. Our passion extends to your entire user experience. We support you every step of the way in your NGS journey and aspirations.

#### STEP 1: Let's find the right solution for you

It's all about your lab's needs now and in the future. We help you determine the right system. Then our in-person trainings and online tools help you discover how to fully expand your research.

#### STEP 2: Setting up

From library prep to informatics, we'll help you achieve operational excellence, with an optimized workflow that will help you run your lab in a cost- and time-efficient manner.

#### **STEP 3: Maintenance and support**

As a global company with 25 years of experience, we not only get you started, we keep your lab running smoothly. We have the infrastructure, teams, and expertise to give you consistent, superior service.

#### Maximum productivity

Illumina Proactive is enhanced service and support you'll come to rely on. Connect your instruments to your free, customized Mylllumina dashboard for instrument analysis and troubleshooting. You'll receive real-time updates on your run progress and instrument utilization. Proactive risk detection by our support team can minimize unplanned downtime and increase sample success.

### You're changing the world. We're right beside you.

Illumina strives to be the best partner possible, providing groundbreaking genomics innovations, ultimate user experience, and exceptional customer service. With a global presence, you'll receive the support you need to facilitate your success. Wherever you are in the world, we provide the talent, resources, and solutions to maximize your discovery power.

Our goal is to apply emerging technologies to the analysis of genetic variation and function, making studies possible that were unimaginable just a few years ago.

And this is precisely the power of Illumina—and you.



Every innovation has led to today—the genome era.

We can't wait to see where we all go next.

### illumına

We are always available for questions, insights, and conversation. Visit us at illumina.com.

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