



Inside the MiSeq[™] sequencing system, you'll find a fully integrated sequencing solution delivering >85% Q30 bases and run times as fast as 8 hours from sample to data.

Integrated approach

• Cluster generation, paired-end fluidics, sequencing, primary and secondary analysis—all on one instrument

Powered by TruSeq[™] for industry-leading data quality

- The same, proven sequencing by synthesis chemistry that's in all Illumina sequencers
- The highest yield of error-free reads and the most bases over Q30
- TruSeq technology has been cited in more than 1,900 publications over the past five years

Fastest time to results

- Total workflow time of just 8 hours
- Direct reporting of variants in 2 hours
- Just 30 minutes hands-on time

Better outside.

Promy to being Cannot Schemer

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Miseq

Outside you'll find a streamlined user experience. Individually tracked load-and-go reagents. Intuitive software interface. Minimal hands-on time.

Efficient operation

- Single-tube Nextera[™] sample preparation in 1.5 hours
- Touch-screen guides you step by step
- Pre-flight check ensures everything is in place for your run
- Keyed flow cell with auto alignment for error-free loading

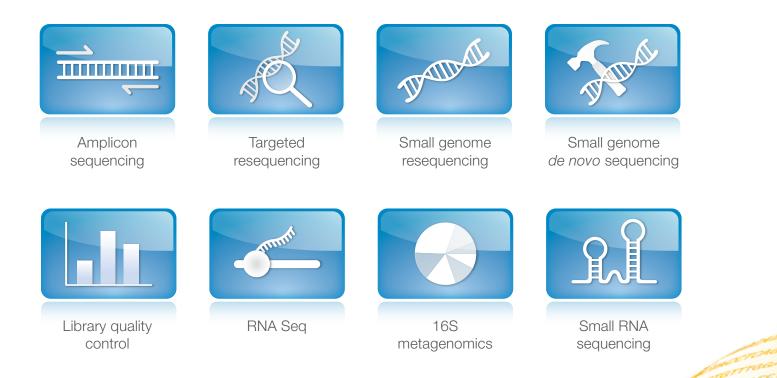
Flexible data management

- Seamlessly perform onboard secondary data analysis with MiSeq Reporter
- BaseSpace[™], our new collaborative cloud-based analysis environment, allows you to store and share data



illumina

The upside is what you can do with it. MiSeq offers built-in scalability. Data compatibility with other Illumina sequencers. The broadest application base.



Side by side with MiSeq.



WORLDWIDE HEADQUARTERS

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Bring more upside to your next sequencing study. Go to www.illumina.com/MiSeq

