

# MiSeq<sup>®</sup> system applications

illumina

# Choose your application. Load, and go.

### Focused power. Speed and simplicity for targeted and small-genome sequencing.

Optimized sample preparation kits, push-button sequencing, and automated data analysis create the first end-to-end sequencing solution. MiSeg delivers a simple workflow and accurate data for access to a wide breadth of applications. And no need for extra hardware-store, analyze, and share data with Illumina's BaseSpace<sup>®</sup> cloud. Sample to data in a single day. All at the touch of a button.



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Amplicon Sequencing

Targeted Resequencing

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# The fastest and easiest sample prep.

### Nextera® XT DNA Sample Prep Kit

Nextera technology combines with the MiSeq system to provide the fastest and easiest solution for amplicon and small-genome sequencing. In one quick step, DNA is simultaneously fragmented and tagged with sequencing adapters to deliver libraries that can be rapidly sequenced with the MiSeq system.

#### Rapid sample preparation

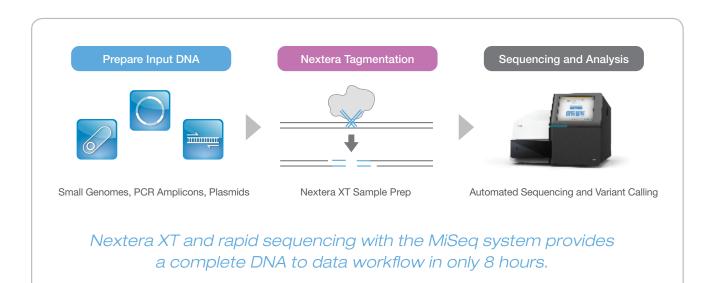
Complete sample prep in as little as 90 minutes with only 15 minutes of hands-on time

*Fastest time to results* Go from DNA to data in only 8 hours

**Optimized for small genomes, PCR amplicons, and plasmids** One sample prep kit for many applications

#### Innovative sample normalization

Eliminates the need for library quantification prior to sample pooling and sequencing



# Amplicon sequencing transformed.

### TruSeq<sup>®</sup> Custom Amplicon Kit

Optimized for the MiSeq system, TruSeq Custom Amplicon (TSCA) offers a highly multiplexed amplicon and sample assay for powerful, fast, and cost-effective variant identification. Requiring just 150 ng of starting gDNA, TSCA is amenable to a wide range of samples, including non-human samples and samples available in limited quantity. Data analysis is performed directly on the MiSeq system, providing quick, easy-to-interpret results for both germline and somatic variation.

#### Simplest assay customization

Easily create projects online with DesignStudio<sup>™</sup> for a range of amplicon sizes and target genomes

#### Fastest sample processing

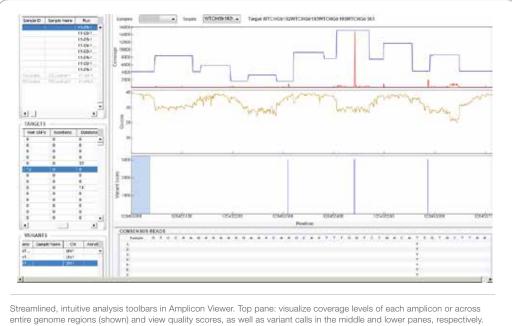
Prepare and enrich up to 1,536 amplicons per reaction and 96 samples per plate simultaneously using standard lab equipment

#### Shortest time to project completion

Go from genomic DNA to fully analyzed data in less than 2 days

#### Easiest analysis

Access pre-configured, on-instrument software for automated variant calling and data analysis



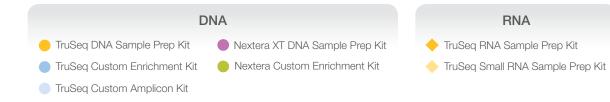
Left scroll bars allow users to navigate through samples and targets, and track variants.

# The widest breadth of sequencing applications.

MiSeq is capable of automated paired-end reads and up to 15 Gb per run, delivering over 600 bases of sequence information. Sample prep kits are optimized for a variety of applications. Empower your research with the MiSeq system.

Application	Recommended Read Length	Illumina Solution
Targeted Resequencing		
Amplicon Sequencing (tens of targets)	1 × 250	•
Amplicon Sequencing (hundreds of targets)	2 × 300	
Hybrid Capture (thousands of targets)	2 × 75	• • •
16S Metagenomics	2 × 150	• •
Clone Checking	1 × 36	•
Small-Genome Sequencing		
De novo	1 × 36, 2 × 300	• •
Resequencing	1 × 36, 2 × 300	• •
Plasmids	1 × 36, 2 × 300	• •
RNA Sequencing		
Small RNA Sequencing	1 × 36	
RNA-Seq (microbial) <sup>†</sup>	2 × 50–250	•
Quality Control		
Library QC	2 × 25	• • • •
Regulation		
ChIP-Seq	1 × 36	•
II Ise with Ribo-Zero™		

<sup>†</sup>Use with Ribo-Zero<sup>™</sup>.



## MiSeq system applications

Focused power. Speed and simplicity for targeted and small-genome sequencing. With the largest applications portfolio at your fingertips. And, access to an entire community of users and tools with BaseSpace. All designed to move your research forward.

The pocket inserts showcase just a few of the applications we've explored with the MiSeq system. And this is just the beginning.

Learn more at www.illumina.com/miseq



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