HiFi sequencing on the Sequel IIe system

HiFi sequencing advantages





Long reads Reads tens of kilobases in length

High accuracy Sequencing free of systematic errors

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Uniform coverage No bias from GC content

Single-molecule resolution Reads from individual native DNA molecules



Epigenetics DNA methylation without special library preparation

Our solution

Sequel[®] lle system

- Supports automated run processing for extended walk-away time
- Provides direct access to HiFi reads
- Delivers 5-base sequencing with methylation status for CpG sites
- Supports the range of HiFi sequencing applications

HiFi consumables

Our consumables support simple, automation-friendly workflows from sample through sequencing:

- Nanobind[®] kits for DNA extraction
- SMRTbell[®] prep kit 3.0 for library preparation
- · Sequel II binding kit 3.2 for sequencing preparation
- Sequel II sequencing kit 2.0 and SMRT® Cell 8M for sequencing







SMRT[®] Link software

Our analytical software tools support you at every step, from run design through analysis



HiFi sequencing applications

Gain comprehensive views of genomes, transcriptomes, and epigenomes.



Whole genome sequencing

Sequence and assemble complete, reference-quality genomes and phase haplotypes



Variant detection

Call all variants — single nucleotide, indel, and structural variants — with high precision and recall

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RNA sequencing

Characterize isoforms with direct sequencing of full-length transcripts no assembly required



Targeted sequencing

Focus on variation in even the most difficult regions of the genome



Metagenomics

Resolve closely related sequences within a heterogeneous mixture



Epigenetics

Directly detect DNA modifications during sequencing

READY TO GET STARTED WITH HIFI SEQUENCING?

Products and services pacb.com/products Documentation pacb.com/documentation Application workflows pacb.com/applications Dublications

Publications pacb.com/pubs





Parts Sequel IIe system (101-986-400) Nanobind CBB big DNA kit (NB-900-001-01) SMRTbell prep kit 3.0 (102-182-700) Binding kit 3.2 and cleanup beads (102-333-300) Sequel II sequencing kit 2.0 (101-820-200)

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Instrument operating environment

| Power requirements | 208–240 VAC, 30A. UPS recommended |
|--------------------------|---|
| Operating temperature | 19–25°C (66.2–77°F) ±2°C per hour |
| Humidity | 20–80%, noncondensing |
| Ventilation | HVAC capacity of up to 13,956 BTU (4090 watts) |
| Nitrogen | 50 psi (2,585 torr) |
| W×D×H | 36.5 × 34 x 66 in (92.7 × 86.4 × 167.6 cm) |
| Weight | 798 lb (362 kg) |

PacBi

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SMRT cell 8M tray (101-389-001)

Technical support 1.877.920.PACB (7222); option 2 support@pacb.com



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