



DeepChek Pipetting Robot Titanium

Automated Liquid Handling Workstation



Providing Lab Efficiency for Optimized Patient Monitoring



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Typical Workflow



UPSTREAM APPLICATIONS

- PCR (DeepChek®-SingleRound Assays, UltraGene® Assays...)
- NGS Library Prep (DeepChek® Library Prep Assays, ...)



DOWNSTREAM APPLICATIONS

- Sequencing
 - Next generation sequencing (NGS)
 - Sanger sequencing
- Genotyping (SNP) fragment analysis
- Cloning

Applications

- Automation for PCR reaction (1 to 96 samples in 96-well plates with any combination of Master Mix, Primers, and Samples)
- Full automation for beads purification

- Automation for Library Preparation for Next-Generation Sequencing
- Normalizing of samples
- Dilution series
- Multiple Mix preparation



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Workflows Overview



LIBRARY PREPARATION







Fragmentation reaction



End-repair & A-tailing plate set-up



End-repair & Atailing reaction



Adaptor ligation set-up & reaction on-deck





SPRI clean-up & PCR mix distribution



PCR

LIBRARY CLEAN-UP



Centrifuge



NORMALIZATION



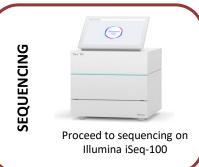
Quantification



Normalization and Pooling in a single tube



Automated protocol is complet





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Key Benefits



REPRODUCIBILITY

Provides run-to-run consistency and reproducibility, thereby enabling verifiable science

EASE OF USE

Automated plate prep is ergonomic and requires minimal intervention from the researchers in the lab.

COMPATIBILITY

Output material is ready to run on a BioAnalyzer® or feed into downstream procedures in your lab or a core facility.

FLEXIBILITY

Pipetting script can be customized to match the required volumes, number of washes in your polymerase chain reaction (PCR) cleanup procedure.

REPRODUCIBILITY

Error-prone, manual steps are automated for greater confidence in performance





- Better reproducibility
- Less manual preparation (more walk-away time)
- Complex, difficult procedures without errors
- Fully automated pipetting
- Automates technically challenging tasks
- 96-well format allows multiple samples to be processed at once
- Customers can walk away during the tedious procedure while maintaining high quality sample integrity



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Workflows Comparison



NGS Library Preparation

for a run of 24 samples

Normalization





■ Handling time: 1min

Robot: 10min

Handling time: 2min

Robot: 30min

Handling time: 1min

Robot: 2min

Pipetting



■ Handling time: 20min

Handling time: 25min

■ Handling time: 10min



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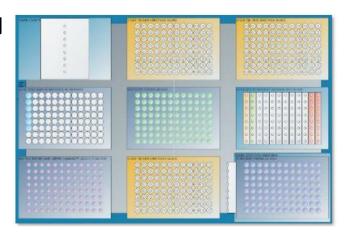
TYPICAL APPLICATION WORKFLOW



PCR PURIFICATION FOR NGS LIBRARY PREPARATION (DeepChek® Library Prep Assay,...)

WALK AWAY AUTOMATION

- Mix beads (in reservoir)
- 2. Add beads to the samples
- 3. Pull down beads
- 4. Remove supernatant
- 5. Wash the beads (80% EtOH)
- 6. Air dry beads to remove EtOH
- 7. Elute sample from beads
- 8. Transfer eluent to new plate



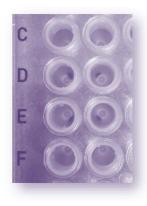
MAGNETIC BEAD RACK

Automatically toggles between disengaged and engaged magnet positions.

 No user intervention is required to pellet or resuspend the magnetic beads during the run.









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Specifications



Communication

USB

Connections

- Three USB host ports and one USB device port
- Two inputs (contact closure, TTL), two relay outputs, and one switched +12V DC 1A output

Control

Touchscreen tablet, laptop, or desktop computer control via USB and TRILUTION® micro software

Dimensions (W x D x H)

- 50.8 x 64.3 x 49.5 cm (20 x 25.3 x 19.5 in)
- Weight: 24.9 kg (55 lbs.)

Volumetric Specification for Transfers

Maximum Permissible Errors

Pipette Head	Volume of DistilledWater	Systematic Error (μL)	Random Error (μL)	Systematic Error (%)	Random Error (%)
MAX8x20	1 μL	±0.08	≤0.05	8	5
	10 μL	±0.15	≤0.10	1.5	1
	20 μL	±0.25	≤0.12	1.25	0.6
MAX8x200	20 μL	±0.50	≤0.16	2.5	0.8
	100 μL	±1.00	≤0.30	1	0.3
	200 μL	±2.00	≤0.50	1	0.25

Positioning Performance

Machine Accuracy (without tips) in XYZ = +/-0.2 mm (0.008")System Accuracy (with current tips) in XYZ = +/-0.9 mm (0.035")



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Product References



DeepChek Pipetting Robot Titanium CE





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