# LUNATIC

## **Genomics Quantification**







#### Get your quant on

Lunatic makes batch quantification of DNA and RNA a no-brainer. All you need is 2  $\mu$ L and 10 minutes to measure up to 96 samples. Load up the tiniest volume and read low concentrations in one step – no standard curves, no dyes. Or take advantage of the wide dynamic range to read high concentrations straight-up, without dilutions. Lunatic gets UV/Vis quantification of nucleic acids on the money every time. Just drop, load and read.



2 μL volume
96 samples
10 minutes
SBS plate format

#### No B.S. workflow

Skip the sample prep, cleaning and worrying about cross-contamination or evaporation – your samples will sit tight for up to 2 hours. Our High Lunatic plate runs 96 samples in 10 minutes. Each microfluidic circuit has two fixed pathlengths built-in to cover a wide dynamic range of 0.03–275 OD. All you do is load up your samples and get kick-ass results.



#### Reliable no matter what

Get spot-on accuracy within 2% and precision within 1%. Using two fixed pathlengths, Lunatic delivers jaw-dropping data at both high and low nucleic acid concentrations. Ditch instruments with moving optics for all the accuracy and precision of fixed UV/Vis pathlengths.



Target Conc. (ng/µL)	Avg Conc. (ng/μL)	Mean Recovery (%)	CV (%)
1400	1395.7	99.7	0.67
700	702.2	100.3	0.28
500	504.4	100.9	0.68
250	250.9	100.4	0.22
140	139.8	99.9	0.41
100	100.5	100.5	0.91
70	68.9	98.4	0.07
50	49.4	98.8	0.67
30	29.9	99.7	0.43
25	24.5	98.0	0.70

#### Always on the money

Don't lose sleep over instrument-to-instrument or lab-to-lab variation. Lunatic loves repeating itself, so standardizing your methods across all labs is easy-peasy. With boatloads of samples at multiple locations, the last thing you need is to struggle with validating the same workflow over and over again.



#### **Measure it all**

Lunatic is the only system out there that can measure nucleic acids at high-throughput and both low and high concentrations. It's got dynamic range that covers from 1.5 to 13,750 ng/ $\mu$ L (dsDNA). On the low end, go for dye-free, quick and easy QC on your precious DNA extract. Run any high concentrated plasmid prep without ever having to dilute again. Get answers in a plate-format and say goodbye to running samples one-by-one.



#### Dilution isn't a solution

Costly fluorescent dyes are only linear for a limited range. At low concentrations Lunatic and PicoGreen agree, while at higher concentrations costly fluorescent dyes saturate. Ditch the time-sucking dye-based workflow, the drag of creating standard curves and the risk of a saturated signal. Know right away if your samples are good to go with Lunatic.



#### Dig up your genomic's dirt

DNA and RNA samples can be freaking messy. Lunatic's unmix applications let you see annoying impurities that other UV/Vis systems would mistake as DNA or RNA. Don't miss the things that can screw up your sequencing – check your preps on Lunatic and know what's in your sample.



#### See plasma clearly

Take the guesswork out of plasma quality determinations. Lunatic eliminates uncertainty by measuring the full UV/Vis spectrum and quantifying the plasma-relevant components: hemoglobin, bilirubin and protein, along with sample turbidity profiles.



#### Set it and forget it

Lunatic also talks to robots. Hook it up with any liquid handler system and run your measurements completely hands-off with our API setup. Stop messing around with instruments that can't be automated and ace your throughput targets.





#### Downstream ready

Smash through the USP & Ph. Eur. UV/Vis requirements for absorbance accuracy, precision, linearity, wavelength accuracy, stray light and resolution with performance verification measurements of independently certified NIST standards. Lunatic is ready when you are to get validated and make the move into QC.



**Performance verification plate** Holds certified pharmacopeia standards for UV/Vis verification



#### Put it on lockdown

GLP labs don't sweat it. Lunatic's software hooks labs up with 21 CFR Part 11 compliant features. We're talking password protection, electronic signatures, full audit trail – the whole package.

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### Specifications

Lunatic instrument specifications			
Dimensions	37 cm W x 46 cm D x 33 cm H; 21 kg		
Operating voltage	24 VDC, 30 W (max)		
Connection	USB, TCP/IP (Service, SiLA)		
Light source	Xenon flash lamp		
Detectors	UV/Vis polychromatic spectrophotometer		
Approval	CE, FCC, CSA		
Wavelength range	230-750 nm		
Wavelength accuracy	≤400 nm: ±1 nm; ≥400 nm: ±2 nm		
Spectral resolution	Better than 2 nm (toluene in hexane)		
Absorbance precision (1 cm quartz cuvette)	<1 OD: ±0.005 OD st dev	1-2 OD: ±0.5% CV	
Absorbance accuracy (1 cm quartz cuvette)	<1 OD: ±0.01 OD	1-2 OD: ±1%	
Lunatic plate specifications	High Lunatic Plate	Lunatic Plate	
Application area	High concentration samples, typically proteins	Low concentration samples, typically nucleic acids	
Application area Sample retention time	High concentration samples, typically proteins Up to 2 hours	Low concentration samples, typically nucleic acids Up to 2 hours	
Application area Sample retention time Recommended sample volume	High concentration samples, typically proteins Up to 2 hours 2 µL	Low concentration samples, typically nucleic acids Up to 2 hours 2 µL	
Application area Sample retention time Recommended sample volume Pathlength(s)	High concentration samples, typically proteinsUp to 2 hours2 μL0.1 mm & 0.7 mm path	Low concentration samples, typically nucleic acids Up to 2 hours 2 µL 0.5 mm path	
Application area Sample retention time Recommended sample volume Pathlength(s) Measurement time for full plate	High concentration samples, typically proteinsUp to 2 hours2 μL0.1 mm & 0.7 mm path~10 minutes	Low concentration samples, typically nucleic acids Up to 2 hours 2 µL 0.5 mm path ~5 minutes	
Application area Sample retention time Recommended sample volume Pathlength(s) Measurement time for full plate Measurement range: OD 10 mm ng/µL dsDNA mg/mL ave protein mix	High concentration samples, typically proteinsUp to 2 hours2 μL0.1 mm & 0.7 mm path~10 minutes0.03-275 OD 10 mm 1.5-13750 ng/μL 0.03-275 mg/mL	Low concentration samples, typically nucleic acids Up to 2 hours 2 µL 0.5 mm path ~5 minutes 0.03-40 OD 10 mm 1.5-2000 ng/µL 0.03-40 mg/mL	
Application area Sample retention time Recommended sample volume Pathlength(s) Measurement time for full plate Measurement range: OD 10 mm ng/µL dsDNA mg/mL ave protein mix Absorbance precision (10 mm pathlength)	High concentration samples, typically proteins Up to 2 hours 2 µL 0.1 mm & 0.7 mm path ~10 minutes 0.03-275 OD 10 mm 1.5-13750 ng/µL 0.03-275 mg/mL <1 OD: ±0.01 OD st dev 1-200 OD: ±1% CV	Low concentration samples, typically nucleic acids Up to 2 hours 2 µL 0.5 mm path ~5 minutes 0.03-40 OD 10 mm 1.5-2000 ng/µL 0.03-40 mg/mL <1 OD: ±0.01 OD st dev >1 OD: ±1.5% CV	
Application area         Sample retention time         Recommended sample volume         Pathlength(s)         Measurement time for full plate         Measurement range:         OD 10 mm         ng/µL dsDNA         mg/mL ave protein mix         Absorbance precision (10 mm pathlength)         Absorbance accuracy (10 mm pathlength)	High concentration samples, typically proteins         Up to 2 hours         2 μL         0.1 mm & 0.7 mm path         ~10 minutes         0.03-275 OD 10 mm         1.5-13750 ng/μL         0.03-275 mg/mL         <1 OD: ±0.01 OD st dev	Low concentration samples, typically nucleic acids Up to 2 hours 2 µL 0.5 mm path ~5 minutes 0.03-40 OD 10 mm 1.5-2000 ng/µL 0.03-40 mg/mL <1 OD: ±0.01 OD st dev >1 OD: ±1.5% CV <1 OD: ±0.02 OD >1 OD: ±4%	
Application area         Sample retention time         Recommended sample volume         Pathlength(s)         Measurement time for full plate         Measurement range:         OD 10 mm         ng/µL dsDNA         mg/mL ave protein mix         Absorbance precision (10 mm pathlength)         Absorbance accuracy (10 mm pathlength)         Dimensions	High concentration samples, typically proteins         Up to 2 hours         2 μL         0.1 mm & 0.7 mm path         ~10 minutes         0.03-275 OD 10 mm         1.5-13750 ng/μL         0.03-275 mg/mL         <1 OD: ±0.01 OD st dev	Low concentration samples, typically nucleic acids Up to 2 hours 2 µL 0.5 mm path ~5 minutes 0.03-40 OD 10 mm 1.5-2000 ng/µL 0.03-40 mg/mL <1 OD: ±0.01 OD st dev >1 OD: ±0.02 OD >1 OD: ±4%	





#### **Unchained Labs**

6870 Koll Center Parkway Pleasanton, CA 94566 Phone: 1.925.587.9800 Toll-free: 1.800.815.6384 Email: info@unchainedlabs.com

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