

Biowave 3 & Biowave 3+ Spectrophotometers Users Manual



biochrom
a division of
Harvard Bioscience, Inc.

TABLE OF CONTENTS

WARRANTY AND REPAIR	5
Warranty Policy	5
SAFETY INFORMATION	6
Hazards and Warnings	6
INTRODUCTION	8
Biochrom WPA Spectrophotometers	8
INSTALLATION	9
Unpacking	9
Positioning	9
Installing	9
INSTRUMENT OVERVIEW	10
Scope	10
Spectrophotometer Principle and Intended Use	10
Hardware	10
Technical Specifications	11
Display and Keypad	11
Instrument Connections	11
PVC PC Software	12
Biochrom Resolution PC Software	12
Instrument Data Output	12
Performing a Measurement	12
USER INTERFACE	13
Keypad	13
Instrument Firmware	13
Home Screen	14
Applications	14
<i>Single Wavelength</i>	14
<i>Concentration</i>	16
<i>Wavescan</i>	18

TABLE OF CONTENTS

<i>Kinetics</i>	19
<i>Standard Curve</i>	21
<i>Multi Wavelength</i>	24
<i>Absorbance Ratio</i>	26
Nucleic Acids	28
<i>DNA</i>	28
<i>RNA</i>	30
<i>Oligo</i>	32
<i>Fluorescent Dye</i>	34
<i>T_m Calculation</i>	37
Protein	39
<i>Protein UV</i>	39
<i>BCA</i>	41
<i>Bradford</i>	44
<i>Lowry</i>	46
<i>Biuret</i>	49
OD 600	51
Favourites, Methods, and USB Memory Stick	53
Settings	55
<i>Date and Time</i>	55
<i>Regional</i>	56
<i>Printer</i>	56
<i>Preferences</i>	56
<i>Contrast</i>	57
<i>About</i>	57
Additional Options	58
Status Bar Icons	60
USEFUL CALCULATIONS	61
Beer-Lambert Law	61
Nucleic Acid Concentrations	62

TABLE OF CONTENTS

Protein Concentrations	63
Nucleic Acid and Protein Purity Ratios	64
Fluorescent Dye Quantity	64
Fluorescent Dye Concentration	64
Fluorescent Frequency of Incorporation (FOI)	65
Fluorescent Dye Incorporation	65
Melting Temperature T_m	65
OD 600	67
TROUBLESHOOTING	68
BUILT-IN PRINTER	69
Built-in Printer Accessory Part Numbers	69
Printer Installation Guide	69
Refilling the Printer Paper	70
ORDERING INFORMATION	71

WARRANTY AND REPAIR

Warranty Policy

Biochrom warrants these instruments for a period of 24 months (2 years), and an additional 12 months (3 years in total) for the xenon lamp, from the date of purchase. from the date of purchase. Where appropriate, Biochrom will repair or replace the unit for defects of workmanship or materials. This warranty does not extend to damage resulting from misuse, neglect, abuse, normal wear and tear, or accidental damage. This warranty extends only to the original purchaser.

IN NO EVENT SHALL BIOCHROM BE LIABLE FOR INCIDENTAL OR CONSEQUENTIAL DAMAGES¹

THERE ARE NO IMPLIED WARRANTIES OF MERCHANTABILITY, FITNESS FOR A PARTICULAR USE, OR OF ANY OTHER NATURE.¹

Biochrom shall not be liable for any claims of any kind whatsoever, as to the equipment. Warranty is void if the instrument is modified, disassembled, repair carried out using unauthorized parts or by a service engineer not previously approved by Biochrom.

Returns

If any defect arises within or outside the warranty period, please contact:

US Office Technical Support

Email: support@hbiosci.com

Telephone (Toll Free, US only): 800-272-2775

Telephone (Outside the US): 508-893-8999

Address:

84 October Hill Road

Holliston, Massachusetts 01746, USA

UK Office Technical Support

Email: support@hbiosci.com

Telephone: +44 (0) 1223 423723

Address:

1020 Cambourne Business Park

Cambourne, Cambridge, UK, CB23 6DW

Goods will not be accepted for return unless the RMA (Return Materials Authorization) number has been issued. The unit must be returned with the completed RMA forms and the Decontamination checklist. The customer is responsible for shipping charges unless the failure is within 30 days of receiving the goods. Please allow a reasonable period of time for completion of repairs or replacement.

Caution Notice

The Biochrom Biowave systems are intended for laboratory use only and can be used in research and development applications. These systems have been designed to meet the standards for electromagnetic compatibility (EMC) and safety intended for laboratory equipment applications. This product should not be used in the presence of a flammable atmosphere such as an anesthetic mixture with air, oxygen, or nitrous oxide.

¹ Where the territory does not allow this exclusion or limitation, this term will not apply.

SAFETY INFORMATION

Hazards and Warnings

This section describes potential hazards which may exist in the operation of these units. A number of warning labels and symbols are affixed to your instrument. These symbols are used to inform you of potential dangers which may exist or where caution is required. Before installing your new unit, please take time to familiarise yourself with these warnings and symbols.

This instrument is subject to the following identified hazards:



This unit uses a Xenon lamp. The lamp energy is mainly confined within the unit but traverses the cell holder when a measurement is being taken. Although the energy present is low and intermittent you are advised not to stare into the beam or attempt to deflect the beam as prolonged exposure could result in permanent eye damage.



High voltages exist within the power supply unit and the Xenon lamp housing. Repair and maintenance should only be carried out by individuals trained to work on these instruments.



There are no biohazardous materials within the unit, however, this unit may be exposed to biohazardous samples during normal laboratory use. To protect users against these hazards we recommend the following decontamination procedures:

- Wipe the exterior casework with disinfectant cleaning wipes.
- Remove cuvettes and cuvette holders.
 - Wash with disinfectant appropriate for the biohazard in question.
 - Rinse with distilled water.
 - Allow to dry thoroughly before reuse.

To further reduce the possibility of biohazards:

- Include an appropriate decontamination certificate for equipment returned for repair.
- Ensure that the operator of the equipment is provided with a safe working environment.

- Use, store and dispose of any chemicals in accordance with manufacturer's guidelines and local safety regulations.
- Provide suitable ventilation when working with volatile solvents or toxic substances.
- Dispose of solvents and chemicals that may be classed as hazardous waste in accordance with local regulatory practice.
- Determine if personal protective equipment (PPE) is required for handling laboratory samples.



All models can be connected to and operated from a PC. To preserve the integrity of the measuring equipment it is essential that the attached PC itself conforms to basic safety and EMC standards and is set up in accordance with the manufacturers' instructions. If in doubt, consult the information that came with your PC.

The following safety precautions should be observed when operating a PC:

- To reduce the chance of eye strain, set up the PC display with the correct viewing position, free from glare and with appropriate brightness and contrast settings.
- To reduce the chance of cross contamination from biological samples, use appropriate personnel protection measures and disinfectant wipes on keyboard and mouse.

In the event of contamination, malfunction or hazard occurring, the operator should disconnect the unit, by removing the power cord, and isolate for decontamination and/or repair.

INTRODUCTION

Biochrom WPA Spectrophotometers

Spectrophotometers are ubiquitous among modern laboratories. Ultraviolet (UV) and Visible (VIS) spectrophotometry has become the method of choice in most laboratories concerned with the identification and quantification of organic and inorganic compounds across a wide range of products and processes. Applied across research, quality, and manufacturing, with continuing focus on life science and pharmaceutical environments, they are equally as relevant in agriculture, animal husbandry and fishery, geological exploration, food safety, environmental monitoring, and many manufacturing industries to name a few.

The WPA Biowave spectrophotometers are quick, accurate, and reliable. They require only small demands on the time and skills of the operator. This operating manual details the processes for taking basic measurements using the Biowave 3 and Biowave 3+ spectrophotometers.

The Biowave 3 instrument is UV-VIS split-beam spectrophotometers with a 5 nm spectral bandwidth and a 10 mm pathlength cell holder. The hardware of the Biowave 3+ model variants differ only by having a narrower 3 nm spectral bandwidth.

INSTALLATION

Unpacking

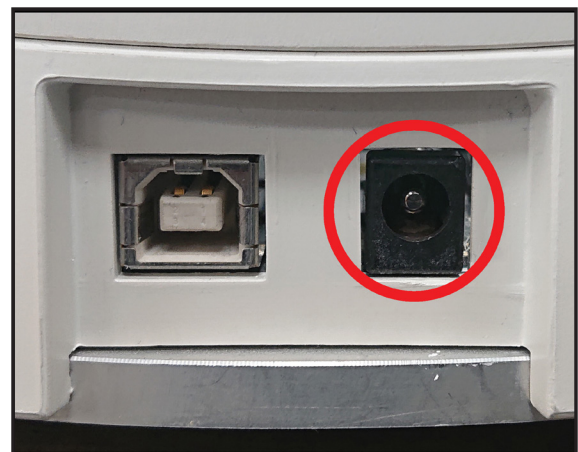
- The unit weighs less than 4 kg. No special handling is required.
- Please keep the original packaging for transport for service or repair as it has been specifically designed to protect the unit from damage during transit.
- Inspect the instrument and its power supply for any signs of damage caused during transit. If any damage is discovered, do not use the instrument and report the problem to your supplier.

Positioning

- Ensure your proposed installation site conforms to the environmental conditions for safe operation:
 - Indoor use
 - 5°C to 40°C
 - Maximum relative humidity 90% up to 31°C decreasing linearly to 50% at 40°C.
 - Extremes of temperature may require recalibration of the unit for optimal performance.
- The instrument must be placed on a stable, level bench or table capable of supporting its weight allowing sufficient space around the instrument for air to circulate freely.
- The instrument should be positioned so that the power supply cable may be readily removed in the event of a hazard or malfunction.
- Locate the instrument in an atmosphere free from dust and corrosive fumes. Use the dust cover to further protect the instrument when not in use.

Installing

- If the instrument has been stored in a cold environment, it should be allowed to come to room temperature before turning it on to avoid compromising the internal calibration procedure.
- The equipment is operated using an 18 VDC power supply adapter unit. Always use the power supply adapter and mains cords supplied with the instrument.
- Mains power requirements are as follows:
 - 100 to 240 VAC~
 - 50 or 60 Hz
 - The UK style mains cord plug has a user replaceable 3A fuse. Replace only with the same rating and type 3A BS1362.
 - The unit maximum power rating is 40 VA.
- Connect the instrument to the mains power using the main power cord and the 18 VDC power supply adapter unit.
- Switch on the instrument after it has been plugged in. The instrument will perform a series of self-diagnostic checks.



18 VDC power supply socket

INSTRUMENT OVERVIEW

Scope

This user manual covers the following range of WPA UV/Visible spectrophotometers:

Part Number	Description
80-3007-32	WPA Biowave 3
80-3007-33	WPA Biowave 3 with Printer
80-3007-34	WPA Biowave 3 with Bluetooth
80-3007-35	WPA Biowave 3 with Printer and Bluetooth
80-3007-37	WPA Biowave 3+
80-3007-38	WPA Biowave 3+ with Printer
80-3007-39	WPA Biowave 3+ with Bluetooth
80-3007-40	WPA Biowave 3+ with Printer and Bluetooth



Spectrophotometer Principle and Intended Use

UV/Visible spectrophotometers measure the transmission of light through a sample. Samples absorb light based on their unique molecular composition. The amount of absorbance is directly proportional to the sample concentration and the pathlength, which is the distance that the light travels through the sample.

UV/Visible spectrophotometers are used in a number of laboratory environments including: life science, clinical, healthcare and industrial laboratories. In a life science laboratory, UV/Visible spectrophotometers are commonly used to measure the concentration of nucleic acids and proteins.

Hardware

Your spectrophotometer is a simple-to-use UV/Visible instrument with a CMOS array detector (1024 pixels). It has no moving parts, which is the basis of the rapid scanning operating system. The look and operation of the Biowave 3 and 3+ are identical; the only difference between them is the bandwidth. Throughout the rest of this manual, the term Biowave 3 will be used to cover both instruments.

INSTRUMENT OVERVIEW

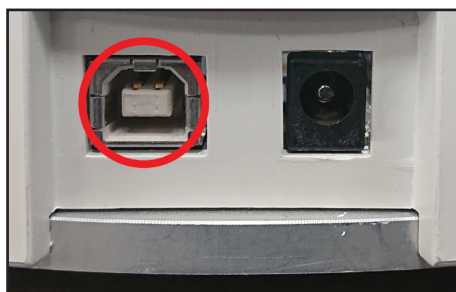
Technical Specifications

Wavelength Range	190 to 1100 nm
Monochromator	Flat grating
Wavelength Calibration	Automatic upon switch on
Beam Height	15 mm
Spectral Bandwidth	5 nm or 3 nm for '+' models
Wavelength Accuracy	±2 nm
Wavelength Reproducibility	±1 nm
Light Sources	Xenon flash lamp
Detector	Twin CMOS array
Photometric Range	-0.300 to 2.500 A, 0.3 to 199 %T
Photometric Linearity	±1.3 % or ±0.008 A whichever is greater at 546 nm
Photometric Reproducibility	±0.002 A to 0.5 A at 546 nm
Stray Light	<0.5 %T 340 nm
Stability	±0.01 A/h at 340 nm
Noise	±0.005 peak to peak ± 0.002 RMS
Digital Output	USB Flash Drive, PC via PVC software, Optional Bluetooth
Data Export	USB Flash Drive: .tsv, native PVC format PC via PVC: .csv, .emf, .xlsx, .xls, .rtf, .tsv, native PVC format
Method Storage	90 with PIN number protection
Sample ID	Yes
Languages	English, German, French, Spanish, Italian, Japanese, Chinese
Dimensions	260 × 390 × 100 mm
Weight	3.00 kg
Power Input	18 VDC at max 40 VA from a supplied 100 to 240 V~, 50/60 Hz Mains Power Adapter

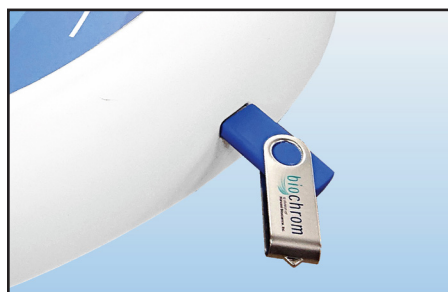
Display and Keypad

The instrument has a ¼ VGA (320 × 240 pixel) resolution backlit LCD display. The instrument's built-in firmware is navigated using the hard-wearing, spill-proof membrane keypad.

Instrument Connections



USB connector for PC connection



USB connector for USB memory stick

INSTRUMENT OVERVIEW

PVC PC Software

The instrument is supplied with the PVC software program (supplied with its own devoted operating manual) on the accompanying USB flash drive. The instrument can be connected to a PC onto which the PVC software has been installed, via a USB A to USB B cable or the factory fit Bluetooth accessory. This enables the operator to “print through” the PC directly to the printer that is connected to it. The data may also be stored as a comma-separated value (.csv), enhanced meta file (.emf), Excel spreadsheet (.xlsx, .xls), rich text format (.rtf), tab-separated value (.tsv) or in a native PVC format file.

Biochrom Resolution PC Software

When connected to a PC the spectrophotometers can be controlled using the Biochrom Resolution PC software packages. Operation using Biochrom Resolution PC software is described in the Resolution user manual or Resolution help file.

Instrument Data Output

A printer accessory is available for the instrument. This may either be supplied preinstalled or as an optional accessory for end-user installation.

Measurement data can also be exported to a USB flash drive via the USB A socket on the side of the instrument, as either a tab-separated value (.tsv) or native PVC format file.

Performing a Measurement

The optical height (z value) of the instrument is 15 mm. The light path is directed from RIGHT to LEFT through the cell chamber.

The cell holder supplied with the instrument accepts standard 10 mm pathlength quartz, glass or plastic cuvettes. When using a cuvette with a pathlength less than 10 mm, ensure the cell is inserted to the far right of the cell holder and secured using an appropriate packing piece.

12 mm test tubes may be used (for cell cultures for example), however they are not recommended as higher quality data is produced by using disposable cuvettes. If used, align the indicator line on 12 mm test tubes in the same direction to ensure reproducible positioning of the tube.

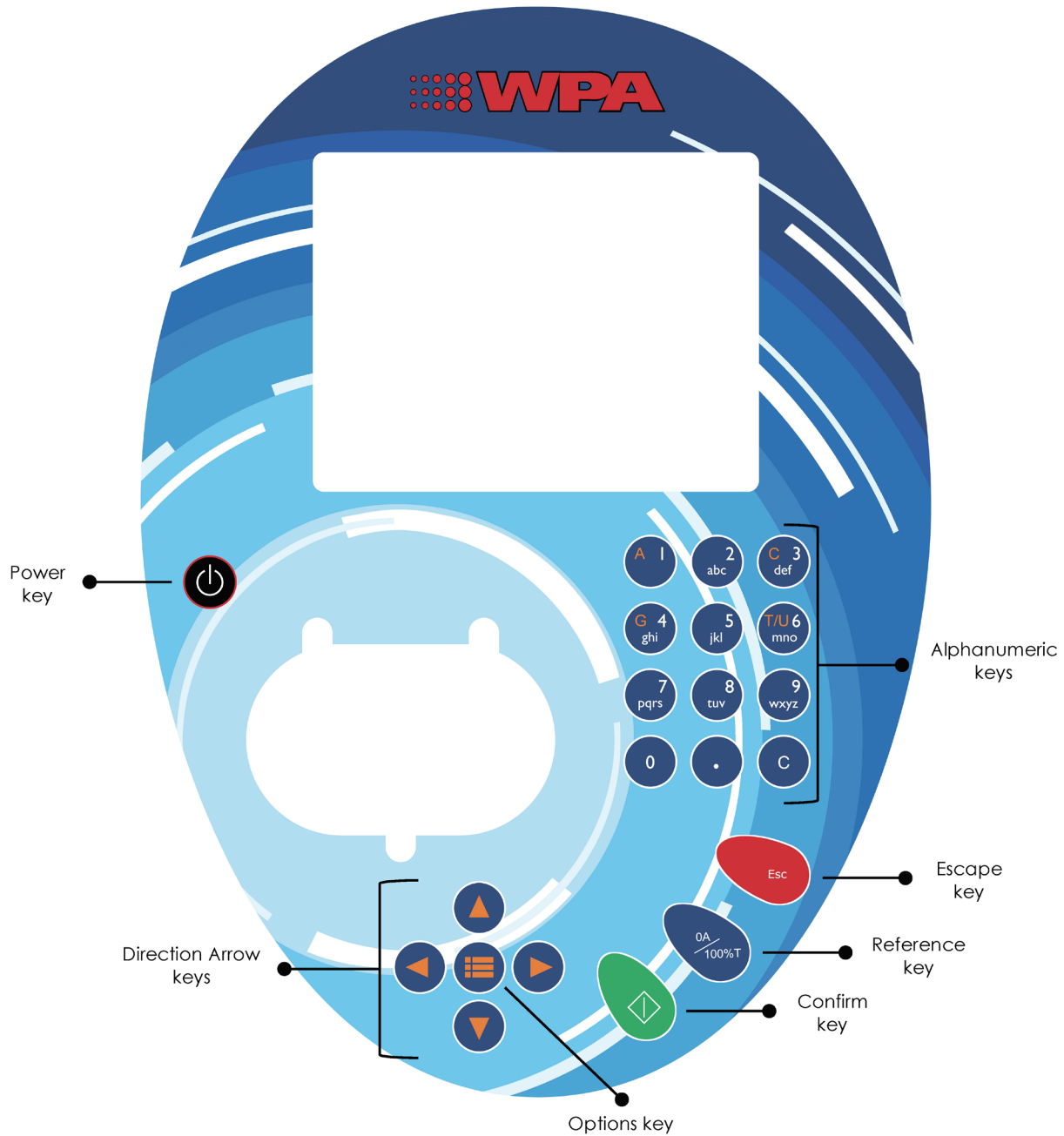
Please consult the “User Interface” section of this user manual for more detail on taking a measurement using the spectrophotometer. In summary, how to perform a measurement is outline below.

1. Open the desired application on the spectrophotometer.
2. Insert a cuvette containing the reference sample into the cuvette holder.
3. Take a reference measurement using the reference key; the acquired reference baseline is applied to any subsequent sample measurements until a new reference baseline is taken, or the application is closed.
4. When the reference is complete, remove the reference sample containing cuvette from the cuvette holder, and replace it with a test sample containing cuvette then take a sample measurement using the confirm key.
5. Repeat step 4 until all the sample data has been collected (see the “Saving and Printing” section for post measurement options).

USER INTERFACE

Keypad

The instrument is controlled using the snap-dome switch-keys of the membrane keypad, offering tactile feedback during operation.



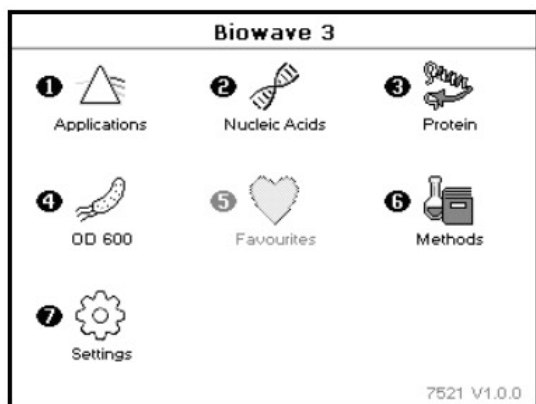
Instrument Firmware

The instrument firmware uses an intuitive menu arrangement that is navigated using the membrane keypad. Upon powering on the instrument, and after completion of the internal calibration, the home screen is displayed.

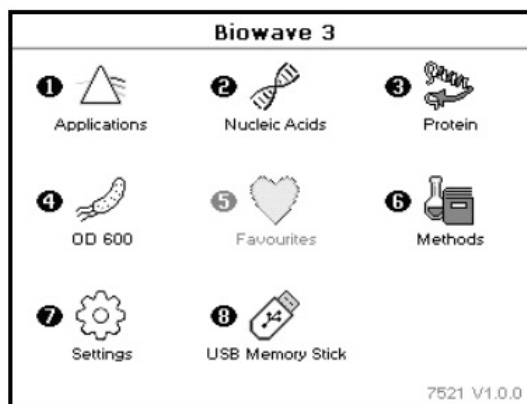
USER INTERFACE

Home Screen

The instrument home screen is the first screen displayed after initialisation of the instrument, and all applications and settings can be accessed from here.



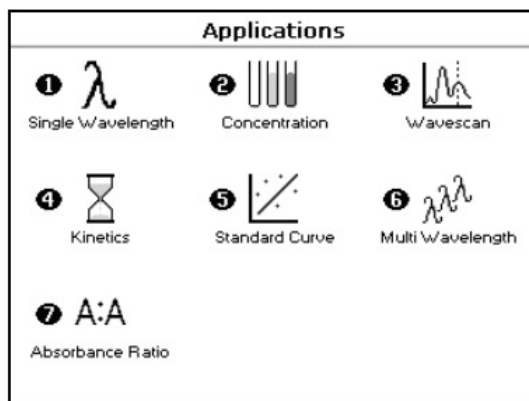
Home screen for the Biowave 3 spectrophotometer



Home screen for the Biowave 3 spectrophotometer displaying the USB memory Stick application, made available when a USB Flash drive is inserted

Applications

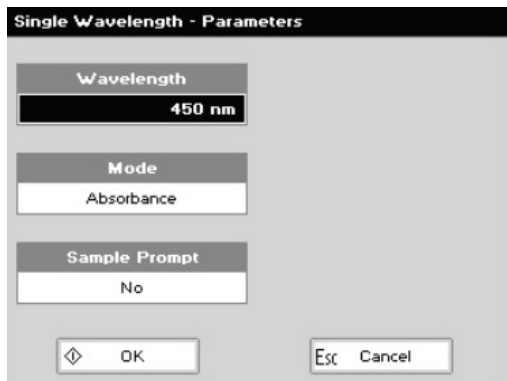
The Applications screen is accessed from the home screen using the '1' key. It contains basic applications with definable parameters to meet the needs of typical laboratory protocols.



USER INTERFACE

Single Wavelength

The Single Wavelength application is accessed from the Applications screen using the '1' key. It can be used to perform simple absorbance (A) or % transmission (%T) measurements.



Step 1

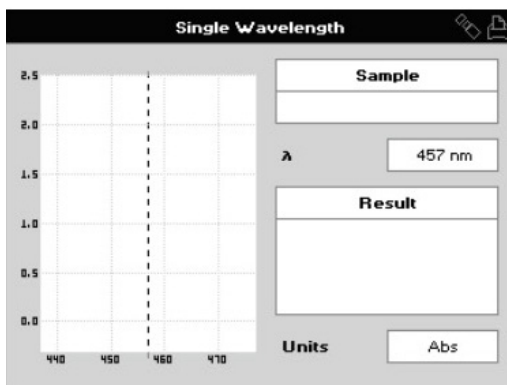
Set the wavelength, 190 to 1100 nm, using the left and right arrow or alphanumeric keys.

Step 2

Press the down arrow, select the mode, "Absorbance" or "%Transmission", using the left and right arrow keys.

Step 3

Press the down arrow, set the sample prompt, "Yes" or "No", to bring up the sample number option before each measurement, using the left and right arrow keys.

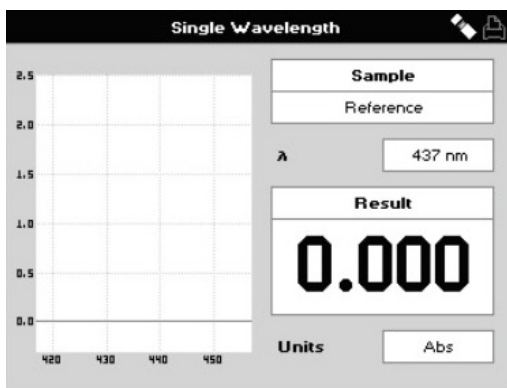


Step 4

Proceed to the measurement screen by selecting "OK" using the confirm key.

OR

Return to the Applications screen by selecting "Cancel" using the escape key.

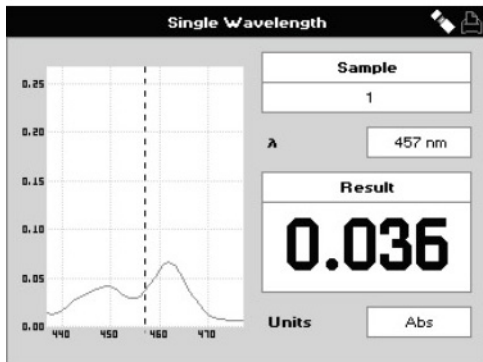


Step 5

Insert the reference sample then take a reference measurement using the reference key.

The acquired reference sample baseline will be applied to all subsequent sample measurements.

USER INTERFACE



Step 6

Replace the previous sample with a test sample then take a sample measurement using the confirm key.

Repeat for all samples.

WARNING!

Previous measurements cannot be recalled so should be printed or output to a USB flash drive before any subsequent samples are measured.

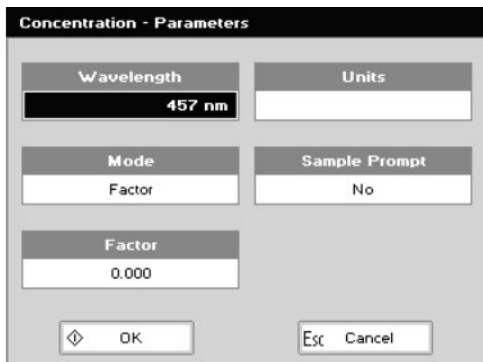
Results

The results are displayed on the screen. Use the left and right arrow keys to move the cursor and display the absorbance at ± 15 nm either side of the set wavelength.

Return to the Applications screen using the escape key OR display the options menu using the options key (see the “Additional Options” section).

Concentration

The Concentration application is accessed from the Applications screen using the ‘2’ key. It can be used to apply a known factor, or one determined using a standard of known concentration to a single wavelength absorbance (A) measurement.



Step 1

Set the wavelength, 190 to 1100 nm, using the left and right arrow or alphanumeric keys.

Step 2

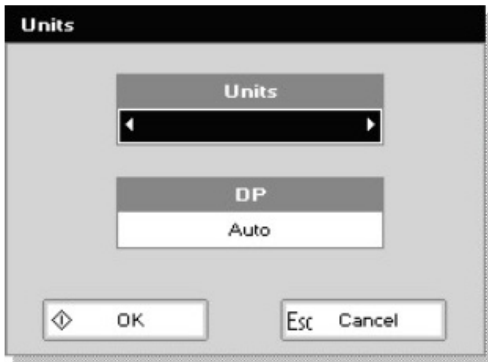
Press the down arrow, select the mode, “Factor”, “Standard” or “Negative Factor”, using the left and right arrow keys.

For “Factor” or “Negative Factor” mode, press the down arrow and enter a value of up to four significant figures, using the alphanumeric keys.

OR

For “Standard” mode, press the down arrow and enter a value of up to four significant figures, for the known concentration of the standard to be used using the alphanumeric keys.

USER INTERFACE



Step 3

Press the down arrow, enter up to 8-digits to define the units that the measurements will be reported in.

OR

Open the Units option using the options key:

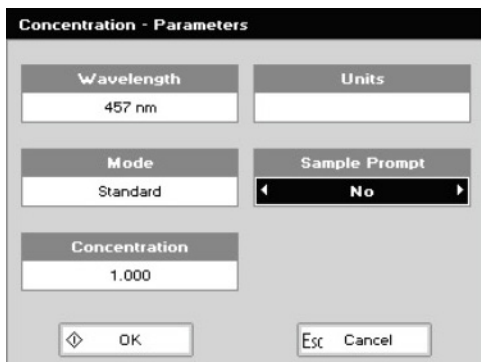
Select one of the predefined units; “ $\mu\text{g/ml}$ ”, “ $\mu\text{g}/\mu\text{l}$ ”, “ $\text{pmol}/\mu\text{l}$ ”, “ mg/dl ”, “ mmol/l ”, “ $\mu\text{mol/l}$ ”, “ g/l ”, “ mg/l ”, “ $\mu\text{g/l}$ ”, “ U/l ”, “%”, “ppm”, “ppb”, or “conc”, using the left and right arrow keys.

Press the down arrow, select the number of decimal places to display the concentration in, “Auto”, “0”, “1”, or “2”, using the left and right arrow keys.

Implement the changes and return to the parameters screen by selecting “OK” using the confirm key.

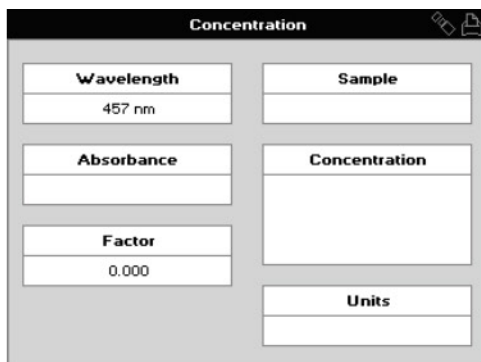
OR

Reject the changes and return to the parameters screen by selecting “Cancel” using the escape key.



Step 4

Press the down arrow, set the sample prompt, “Yes” or “No”, to bring up the sample number option before each measurement, using the left and right arrow keys.



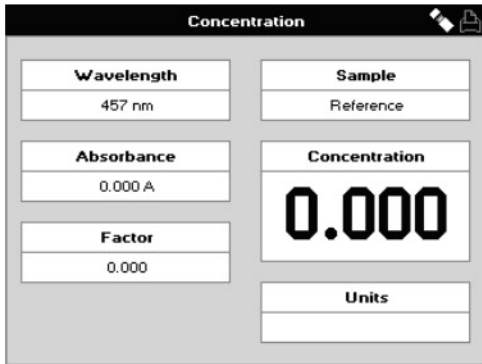
Step 5

Proceed to the measurement screen by selecting “OK” using the confirm key.

OR

“Return to the Applications screen by selecting “Cancel” using the escape key.

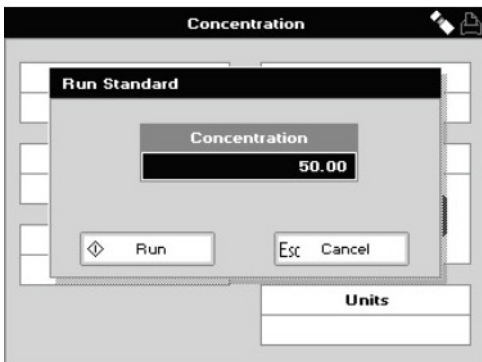
USER INTERFACE



Step 6

Insert the reference sample then take a reference measurement using the reference key.

The acquired reference sample baseline will be applied to all subsequent sample measurements.



Step 7

For "Standard" mode, replace the previous sample with a standard sample then take a standard measurement using the confirm key.

Confirm the concentration of the standard sample by selecting "OK" using the confirm key.

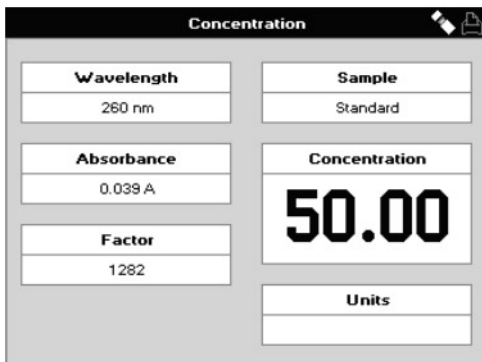
OR

Change the concentration of the standard sample by entering a value of up to four significant figures, using the alphanumeric keys.

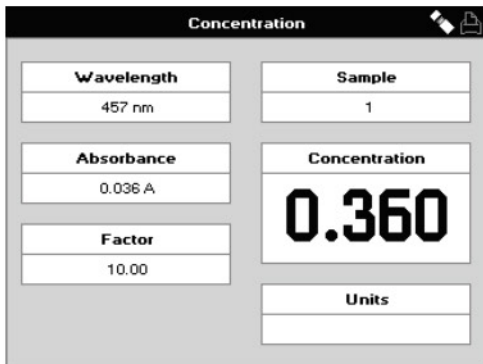
Implement the change and return to the measurement screen to run the standard measurement by selecting "OK" using the confirm key.

OR

Reject the changes, return to the measurement and cancel the standard measurement run by selecting "Cancel" using the escape key.



USER INTERFACE



Step 8

Replace the previous sample with a test sample then take a sample measurement using the confirm key.

Repeat for all samples.

WARNING!

Previous measurements cannot be recalled so should be printed or output to a USB flash drive before any subsequent samples are measured.

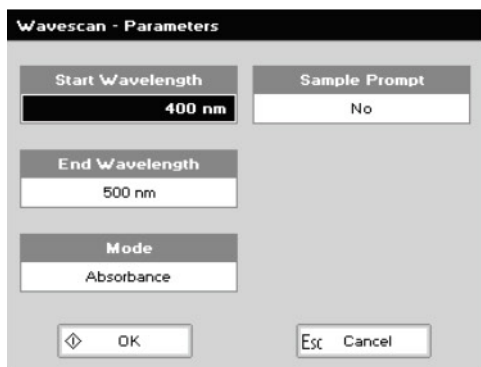
Results

The results are displayed on the screen.

Return to the Applications screen using the escape key OR display the options menu using the options key (see the “Additional Options” section).

Wavescan

The Wavescan application is accessed from the Applications screen using the ‘3’ key. It can be used to perform absorbance (A) or % transmission (%T) measurements across a range of wavelengths creating an absorbance, or transmission, spectrum.



Step 1

Set the start wavelength, 200 to 940 nm, using the left and right arrow or alphanumeric keys.

Step 2

Press the down arrow, set the end wavelength, 210 to 950 nm, using the left and right arrow or alphanumeric keys.

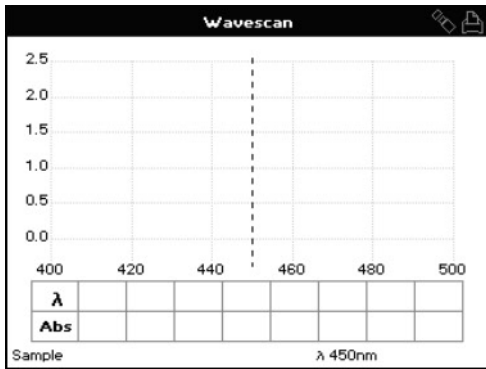
Step 3

Press the down arrow, select the mode, “Absorbance” or “%Transmission”, using the left and right arrow keys.

Step 4

Press the down arrow, set the sample prompt, “Yes” or “No”, to bring up the sample number option before each measurement, using the left and right arrow keys.

USER INTERFACE

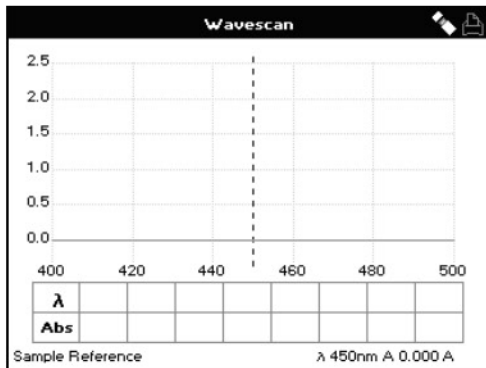


Step 5

Proceed to the measurement screen by selecting “OK” using the confirm key.

OR

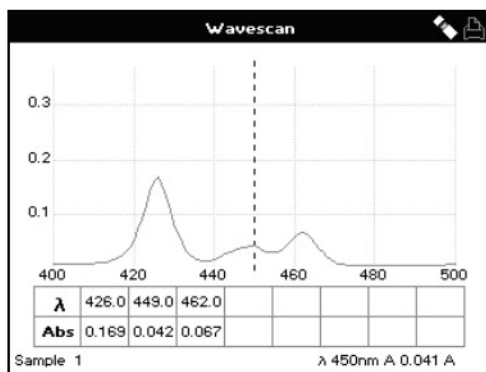
Return to the Applications screen by selecting “Cancel” using the escape key.



Step 6

Insert the reference sample then take a reference measurement using the reference key.

The acquired reference sample baseline will be applied to all subsequent sample measurements.



Step 7

Replace the previous sample with a test sample then take a sample measurement using the confirm key.

Repeat for all samples.

WARNING!

Previous measurements cannot be recalled so should be printed or output to a USB flash drive before any subsequent samples are measured.

Results

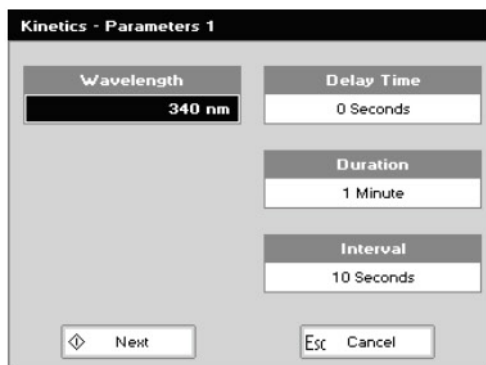
The results are displayed on the screen. Use the left and right arrow keys to move the cursor and display the absorbance at each wavelength position.

Return to the Applications screen using the escape key OR display the options menu using the options key (see the “Additional Options” section).

USER INTERFACE

Kinetics

The Kinetics application is accessed from the Applications screen using the '4' key. It can be used to perform a series of absorbance (A) measurements over a defined timeframe creating a time-course trace.



Step 1

Set the wavelength, 190 to 1100 nm, using the left and right arrow or alphanumeric keys.

Step 2

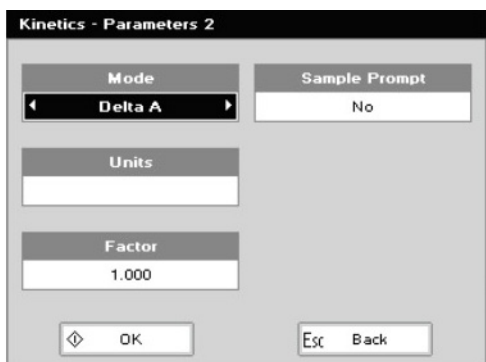
Press the down arrow, set the delay time before the first measurement, 0 to 600 seconds, using the left and right arrow or alphanumeric keys.

Step 3

Press the down arrow, set the duration of the observation, 1 to 60 minutes, using the left and right arrow or alphanumeric keys.

Step 4

Press the down arrow, set the interval between individual measurements, 5 to 60 seconds, using the left and right arrow or alphanumeric keys.



Step 5

Proceed to the second parameters screen by selecting "Next" using the confirm key.

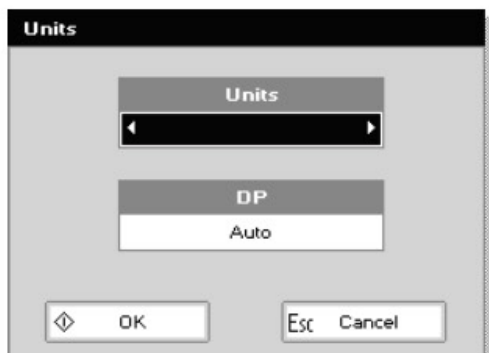
OR

Return to the Applications screen by selecting "Cancel" using the escape key.

Step 6

Select the mode to define the desired final result, "Delta A", "Final A", or "Slope", using the left and right arrow keys.

USER INTERFACE



Step 7

Press the down arrow, enter up to 8-digits to define the units that the final result will be reported in.

OR

Open the units options using the options key:

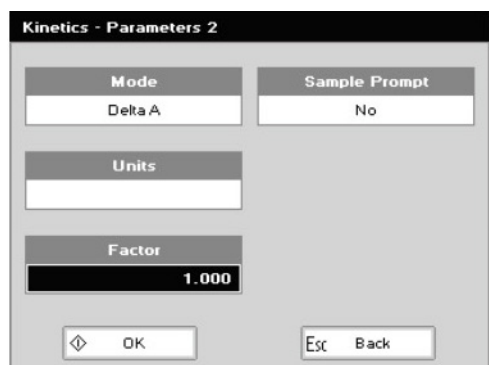
Select one of the predefined units; “ $\mu\text{g}/\text{ml}$ ”, “ $\mu\text{g}/\mu\text{l}$ ”, “ $\text{pmol}/\mu\text{l}$ ”, “ mg/dl ”, “ mmol/l ”, “ $\mu\text{mol}/\text{l}$ ”, “ g/l ”, “ mg/l ”, “ $\mu\text{g}/\text{l}$ ”, “ U/l ”, “%”, “ppm”, “ppb”, or “conc”, using the left and right arrow keys.

Press the down arrow, select the number of decimal places to display the concentration in, “Auto”, “0”, “1”, or “2”, using the left and right arrow keys.

Implement the changes and return to the parameters screen by selecting “OK” using the confirm key.

OR

Reject the changes and return to the parameters screen by selecting “Cancel” using the escape key.

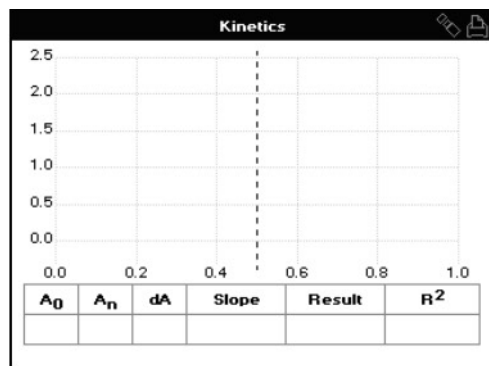


Step 8

Press the down arrow, enter a factor to be used to generate the final result of a value of up to four significant figures, using the alphanumeric keys.

Step 9

Press the down arrow, set the sample prompt, “Yes” or “No”, to bring up the sample number option before each measurement, using the left and right arrow keys.



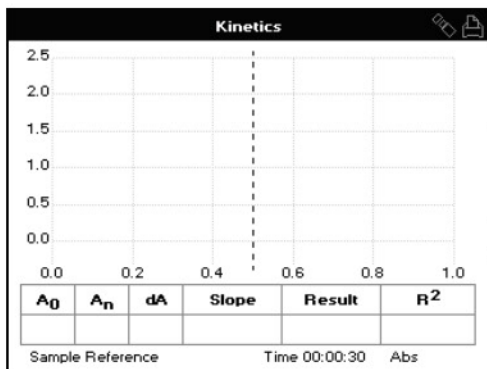
Step 10

Proceed to the measurement screen by selecting “OK” using the confirm key.

OR

Return to the previous parameters screen by selecting “Back” using the escape key.

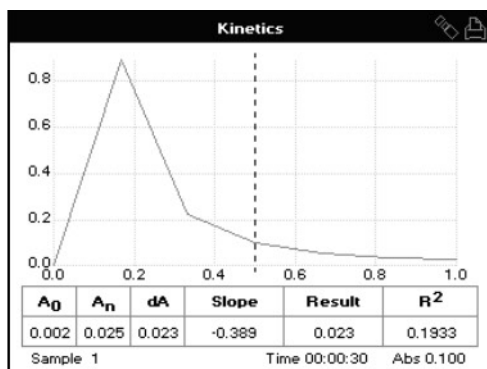
USER INTERFACE



Step 11

Insert the reference sample then take a reference measurement using the reference key.

The acquired reference sample baseline will be applied to all subsequent sample measurements.



Step 12

Replace the previous sample with a test sample then take a sample measurement using the confirm key.

Repeat for all samples.

WARNING!

Previous measurements cannot be recalled so should be printed or output to a USB flash drive before any subsequent samples are measured.

Results

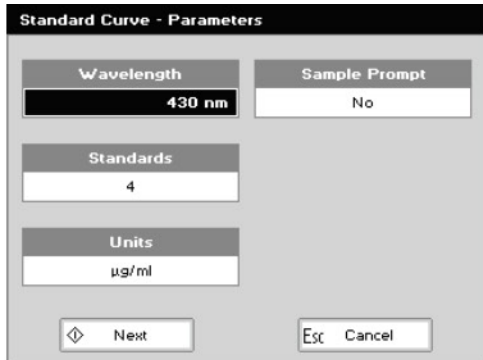
The results are displayed on the screen. Use the left and right arrow keys to move the cursor and display the absorbance at each interval position.

Return to the Applications screen using the escape key OR display the options menu using the options key (see the “Additional Options” section).

Standard Curve

The Standard Curve application is accessed from the Applications screen using the ‘5’ key. It can be used to create a calibration curve from standard samples of known concentration. The curve fit equation is then applied to the absorbance (A) measurements of any subsequent test samples to determine their concentration.

USER INTERFACE

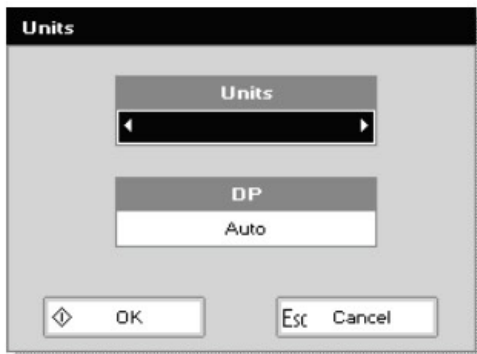


Step 1

Set the wavelength, 190 to 1100 nm, using the left and right arrow or alphanumeric keys.

Step 2

Press the down arrow, set the number of standard samples of known concentration, 1 to 9, using the left and right arrow or alphanumeric keys.



Step 3

Press the down arrow, enter up to 8-digits to define the units that the primary result will be reported in.

OR

Open the units options using the options key:

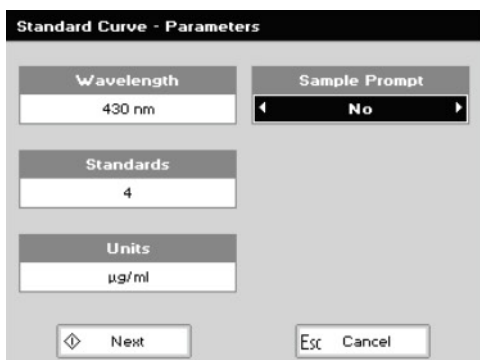
Select one of the predefined units; “µg/ml”, “µg/µl”, “pmol/µl”, “mg/dl”, “mmol/l”, “µmol/l”, “g/l”, “mg/l”, “µg/l”, “U/l”, “%”, “ppm”, “ppb”, or “conc”, using the left and right arrow keys.

Press the down arrow, select the number of decimal places to display the concentration in, “Auto”, “0”, “1”, or “2”, using the left and right arrow keys.

Implement the changes and return to the parameters screen by selecting “OK” using the confirm key.

OR

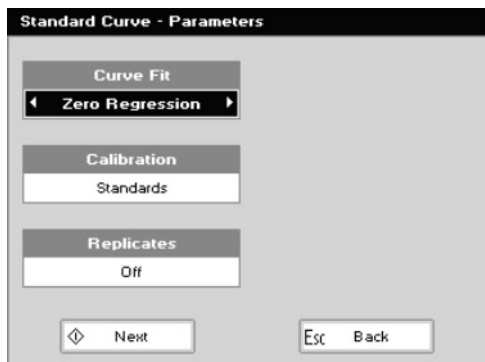
Reject the changes and return to the parameters screen by selecting “Cancel” using the escape key.



Step 4

Press the down arrow, set the sample prompt, “Yes” or “No”, to bring up the sample number option before each measurement, using the left and right arrow keys.

USER INTERFACE



Step 5

Proceed to the second parameters screen by selecting “Next” using the confirm key.

OR

Return to the Applications screen by selecting “Cancel” using the escape key.

Step 6

Select the curve fit, “Zero Regression”, “2nd Order Polynomial”, “Regression”, “Interpolation”, or “Cubic Spline”, using the left and right arrow keys.

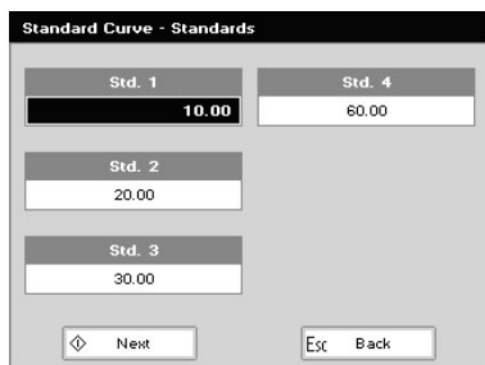
PLEASE NOTE

If only 1 standard sample of known concentration is used to create the standard curve, only a “Zero Regression” fit can be selected. If 2 standard samples are being used to create the standard curve, only “Zero Regression”, “Regression”, and “Interpolation” fits can be selected.

Step 7

Press the down arrow, select the source of the calibration, “Standards”, “Manual”, or “New Standards”, using the left and right arrow keys.

For “Standards” and “New Standards” calibration, press the down arrow and select the number of standard sample replicates, “Off”, “2”, or “3”, using the left and right arrow keys.



Step 8

Proceed to the Standards screen by selecting “Next” using the confirm key.

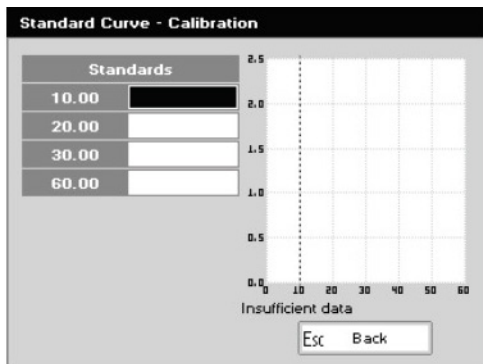
OR

Return to the previous parameters screen by selecting “Back” using the escape key.

Step 9

Enter a value of up to four significant figures, for the known concentration of each standard to be used using the alphanumeric keys.

USER INTERFACE



Step 10

Proceed to the Calibration screen by selecting “Next” using the confirm key.

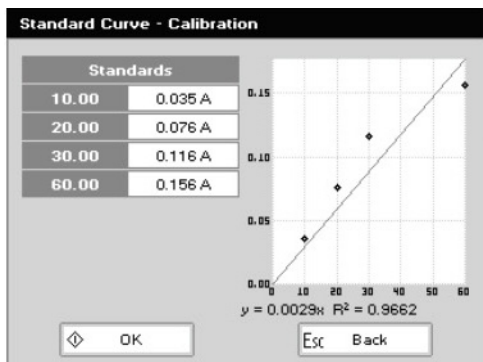
OR

Return to the previous parameters screen by selecting “Back” using the escape key.

Step 11

Insert the reference sample then take a reference measurement using the reference key.

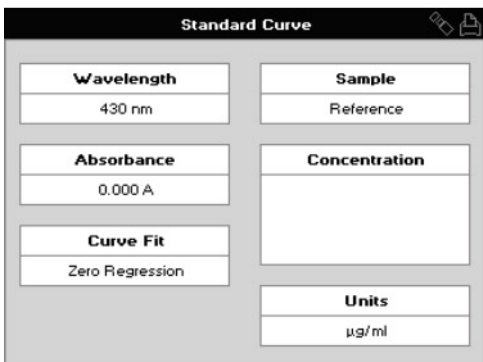
The acquired reference sample baseline will be applied to all subsequent standard and test sample measurements.



Step 12

Replace the previous sample with the first standard sample then take a sample measurement using the confirm key.

Repeat for all standards samples.



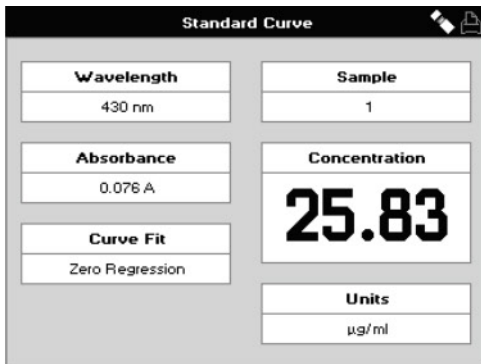
Step 13

Proceed to the measurement screen by selecting “OK” using the confirm key.

OR

Return to the standards screen by selecting “Back” using the escape key.

USER INTERFACE



Step 14

Insert a test sample then take a sample measurement using the confirm key.

Repeat for all samples.

WARNING!

Previous measurements cannot be recalled so should be printed or output to a USB flash drive before any subsequent samples are measured.

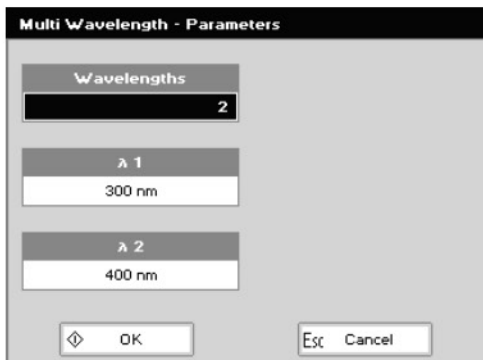
Results

The results are displayed on the screen.

Return to the Applications screen using the escape key OR display the options menu using the options key (see the “Additional Options” section).

Multi Wavelength

The Multi Wavelength application is accessed from the Applications screen using the ‘6’ key. It can be used to perform simple absorbance (A) measurements at up to five specific wavelengths.



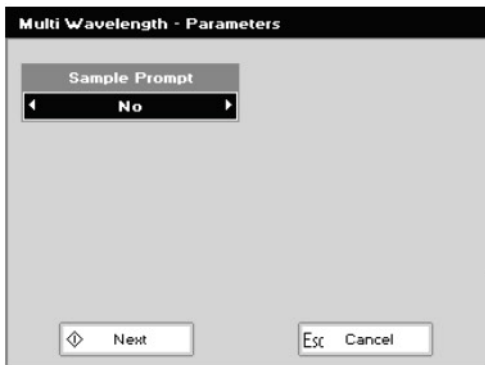
Step 1

Set the number of wavelengths, 2 to 5, using the left and right arrow or alphanumeric keys.

Step 2

Press the down arrow, set the first wavelength, 190 to 1100 nm, using the left and right arrow or alphanumeric keys.

USER INTERFACE



Step 3

Proceed to the second parameters screen by selecting “OK” using the confirm key.

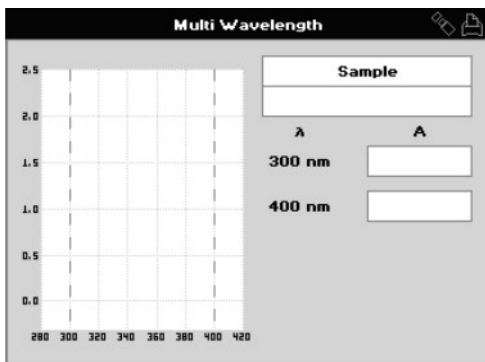
OR

Return to the Applications screen by selecting “Cancel” using the escape key.

Step 4

Press the down arrow, set the sample prompt, “Yes” or “No”, to bring up the sample number option before each measurement, using the left and right arrow keys.

Repeat for all wavelengths.

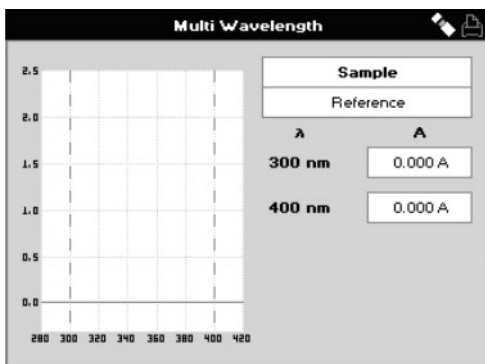


Step 5

Proceed to the measurement screen by selecting “OK” using the confirm key.

OR

Return to the previous parameters screen by selecting “Cancel” using the escape key.

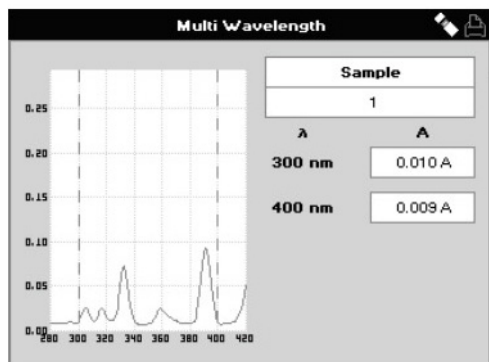


Step 6

Insert the reference sample then take a reference measurement using the reference key.

The acquired reference sample baseline will be applied to all subsequent sample measurements.

USER INTERFACE



Step 7

Replace the previous sample with a test sample then take a sample measurement using the confirm key.

Repeat for all samples.

WARNING!

Previous measurements cannot be recalled so should be printed or output to a USB flash drive before any subsequent samples are measured.

Results

The results are displayed on the screen.

Return to the Applications screen using the escape key OR display the options menu using the options key (see the “Additional Options” section).

Absorbance Ratio

The Absorbance Ratio application is accessed from the Applications screen using the ‘7’ key. It can be used to perform two absorbance (A) measurements and calculate the absorbance ratio between them.

Step 1

Set wavelength 1, 190 to 1100 nm, using the left and right arrow or alphanumeric keys.

Press the down arrow, repeat for wavelength 2.

Step 2

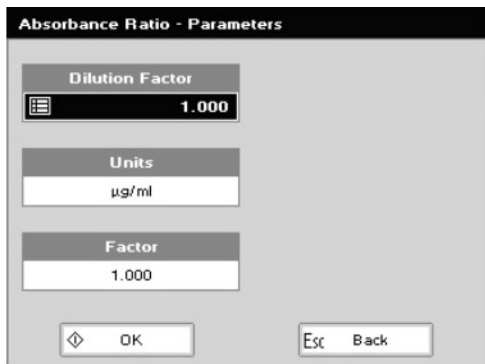
Press the down arrow, set the sample prompt, “Yes” or “No”, to bring up the sample number option before each measurement, using the left and right arrow keys.

Step 3

Press the down arrow, set the background, “On” or “Off”, using the left and right arrow keys.

For background set to “On”, press the down arrow and set wavelength 3, 190 to 1100 nm, using the left and right arrow or alphanumeric keys.

USER INTERFACE

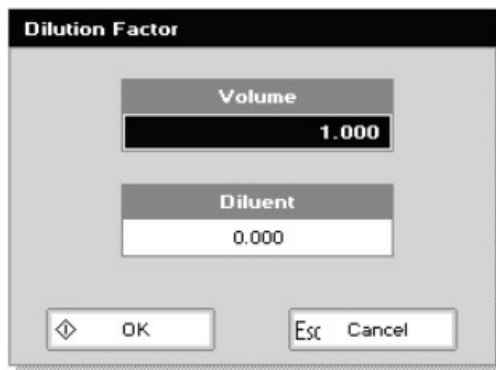


Step 4

Proceed to the parameter screen by selecting “OK” using the confirm key.

OR

Return to the Applications screen by selecting “Cancel” using the escape key.



Step 5

Enter any dilution factor to be applied to the absorbance measurement of a value of up to four significant figures, using the alphanumeric keys.

OR

Open the dilution factor options using the options key:

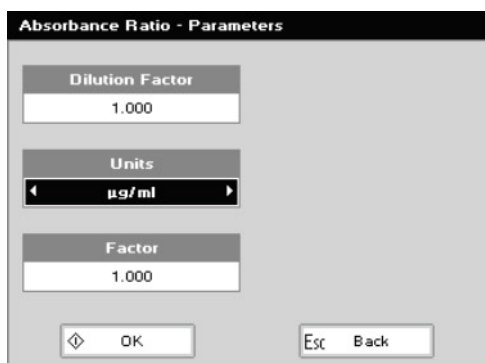
Set the initial sample volume of a value of up to four significant figures, using the alphanumeric keys.

Press the down arrow, set the amount of diluent added to the initial volume of a value of up to four significant figures, using the alphanumeric keys.

Implement the changes and return to the parameters screen by selecting “OK” using the confirm key.

OR

Reject the changes and return to the parameters screen by selecting “Cancel” using the escape key.



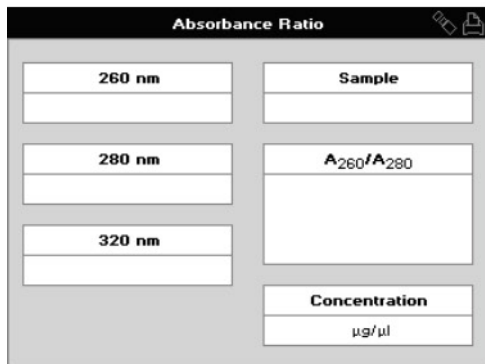
Step 6

Press the down arrow, select one of the predefined units, “µg/ml”, “ng/µl”, or “µg/µl”, using the left and right arrow keys.

Step 7

Press the down arrow, enter a concentration factor of up to four significant figures, using the alphanumeric keys.

USER INTERFACE

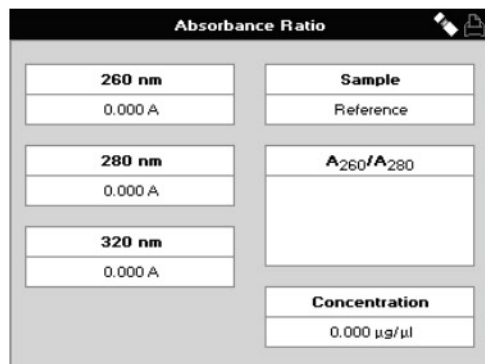


Step 8

Proceed to the measurement screen by selecting “OK” using the confirm key.

OR

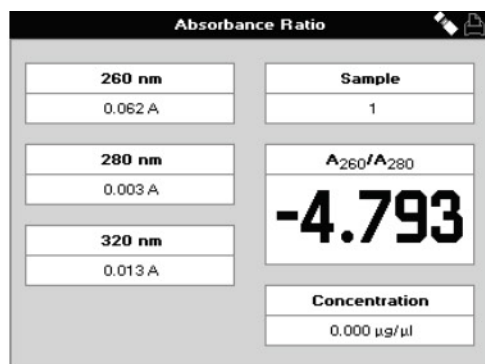
Return to the wavelengths screen by selecting “Back” using the escape key.



Step 9

Insert the reference sample then take a reference measurement using the reference key.

The acquired reference sample baseline will be applied to all subsequent sample measurements.



Step 10

Replace the previous sample with a test sample then take a sample measurement using the confirm key.

Repeat for all samples.

WARNING!

Previous measurements cannot be recalled so should be printed or output to a USB flash drive before any subsequent samples are measured.

Results

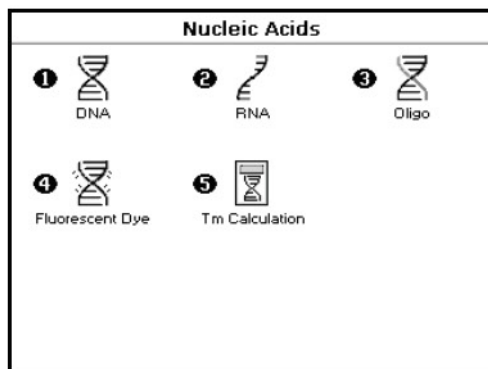
The results are displayed on the screen.

Return to the Applications screen using the escape key OR display the options menu using the options key (see the “Additional Options” section).

USER INTERFACE

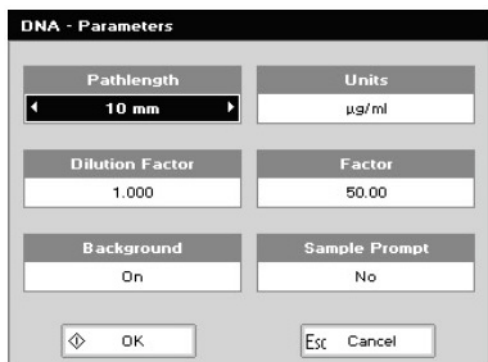
Nucleic Acids

The Nucleic Acids screen is accessed from the home screen using the '2' key. It contains predefined nucleic acid quantification methods and a theoretical melting temperature (T_m) calculation tool. All calculation applied within the Nucleic Acids applications are described in the Useful Calculation section.



DNA

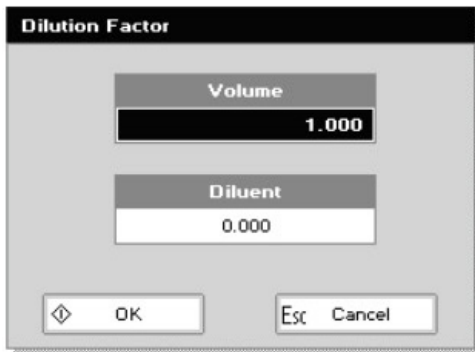
The DNA application is accessed from the Nucleic Acids screen using the '1' key. It can be used to perform DNA quantification measurements at 260 nm and to assess sample quality using A₂₆₀/A₂₈₀ and A₂₆₀/A₂₃₀ absorbance (A) ratios.



Step 1

Set the pathlength; "10 mm", "5 mm", "1 mm", "0.5 mm", "0.2 mm", or "0.125 mm", using the left and right arrow keys.

USER INTERFACE



Step 2

Press the down arrow, enter any dilution factor to be applied to the absorbance measurement of a value of up to four significant figures, using the alphanumeric keys.

OR

Open the dilution factor options using the options key:

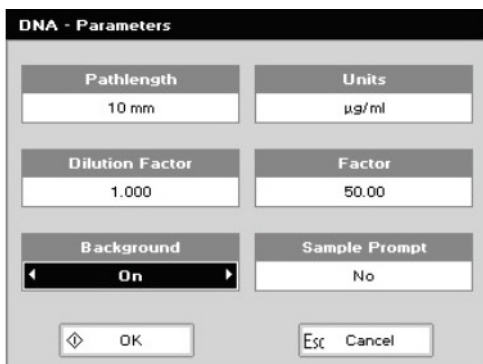
Set the initial sample volume of a value of up to four significant figures, using the alphanumeric keys.

Press the down arrow, set the amount of diluent added to the initial volume of a value of up to four significant figures, using the alphanumeric keys.

Implement the changes and return to the parameters screen by selecting "OK" using the confirm key.

OR

Reject the changes and return to the parameters screen by selecting "Cancel" using the escape key.



Step 3

Press the down arrow, set the background, "On" or "Off", using the left and right arrow keys.

Step 4

Press the down arrow, select one of the predefined units, "µg/ml", "ng/µl", or "µg/µl", using the left and right arrow keys.

For each predefined unit a default factor is applied, "50.00" for "µg/ml" and "ng/µl", or "0.050" for "µg/µl".

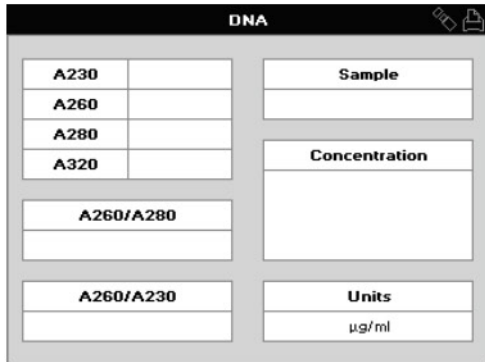
OR

Press the down arrow and enter a custom value of up to four significant figures, using the alphanumeric keys.

Step 5

Press the down arrow, set the sample prompt, "Yes" or "No", to bring up the sample number option before each measurement, using the left and right arrow keys.

USER INTERFACE

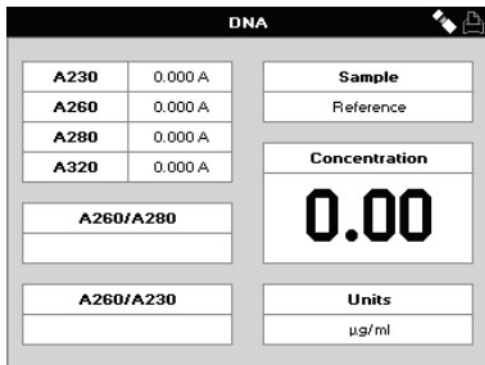


Step 6

Proceed to the measurement screen by selecting “OK” using the confirm key.

OR

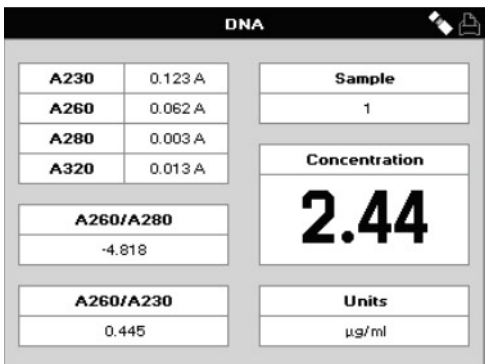
Return to the Applications screen by selecting “Cancel” using the escape key.



Step 7

Insert the reference sample then take a reference measurement using the reference key.

The acquired reference sample baseline will be applied to all subsequent sample measurements.



Step 8

Replace the previous sample with a test sample then take a sample measurement using the confirm key.

Repeat for all samples.

WARNING!

Previous measurements cannot be recalled so should be printed or output to a USB flash drive before any subsequent samples are measured.

Results

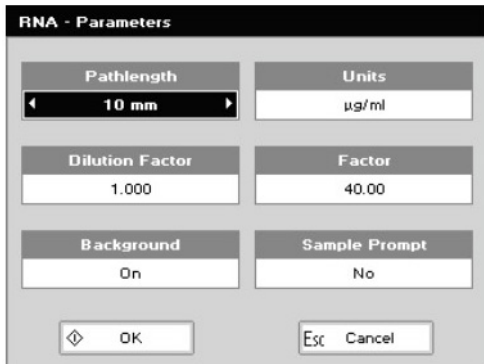
The results are displayed on the screen.

Return to the Applications screen using the escape key OR display the options menu using the options key (see the Additional Options section).

USER INTERFACE

RNA

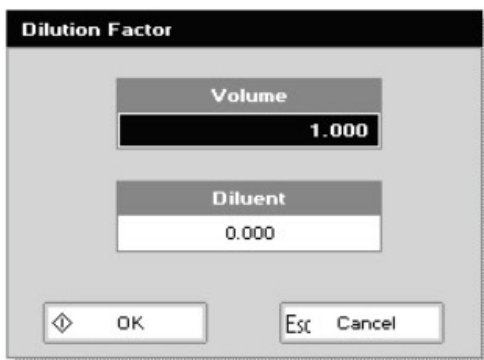
The RNA application is accessed from the Nucleic Acids screen using the '2' key. It can be used to perform RNA quantification measurements at 260 nm and to assess sample quality using A260/A280 and A260/A230 absorbance (A) ratios.



Step 1

Set the pathlength; "10 mm", "5 mm", "1 mm", "0.5 mm", "0.2 mm", or "0.125 mm", using the left and right arrow keys.

Set the pathlength; "10 mm", "5 mm", "1 mm", "0.5 mm", "0.2 mm", or "0.125 mm", using the left and right arrow keys.



Step 2

Press the down arrow, enter any dilution factor to be applied to the absorbance measurement of a value of up to four significant figures, using the alphanumeric keys.

OR

Open the dilution factor options using the options key:

Set the initial sample volume of a value of up to four significant figures, using the alphanumeric keys.

Press the down arrow, set the amount of diluent added to the initial volume of a value of up to four significant figures, using the alphanumeric keys.

Implement the changes and return to the parameters screen by selecting "OK" using the confirm key.

OR

Reject the changes and return to the parameters screen by selecting "Cancel" using the escape key.

USER INTERFACE

RNA - Parameters	
Pathlength	Units
10 mm	µg/ml
Dilution Factor	Factor
1.000	40.00
Background	Sample Prompt
On	No
OK	Esc Cancel

Step 3

Press the down arrow, set the background, “On” or “Off”, using the left and right arrow keys.

Step 4

Press the down arrow, select one of the predefined units, “µg/ml”, “ng/µl”, or “µg/µl”, using the left and right arrow keys.

For each predefined unit a default factor is applied, “40.00” for “µg/ml” and “ng/µl”, or “0.040” for “µg/µl”.

OR

Press the down arrow and enter a custom value of up to four significant figures, using the alphanumeric keys.

Step 5

Press the down arrow, set the sample prompt, “Yes” or “No”, to bring up the sample number option before each measurement, using the left and right arrow keys.

RNA	
A230	
A260	
A280	
A320	
A260/A280	
A260/A230	
Sample	
Concentration	
Units	µg/ml

Step 6

Proceed to the measurement screen by selecting “OK” using the confirm key.

OR

Return to the Applications screen by selecting “Cancel” using the escape key.

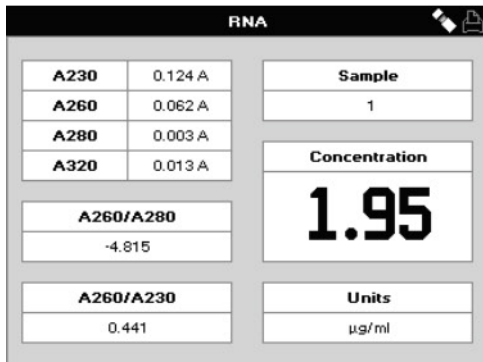
RNA	
A230	0.000 A
A260	0.000 A
A280	0.000 A
A320	0.000 A
A260/A280	
A260/A230	
Sample	Reference
Concentration	0.00
Units	µg/ml

Step 7

Insert the reference sample then take a reference measurement using the reference key.

The acquired reference sample baseline will be applied to all subsequent sample measurements.

USER INTERFACE



Step 8

Replace the previous sample with a test sample then take a sample measurement using the confirm key.

Repeat for all samples.

WARNING!

Previous measurements cannot be recalled so should be printed or output to a USB flash drive before any subsequent samples are measured.

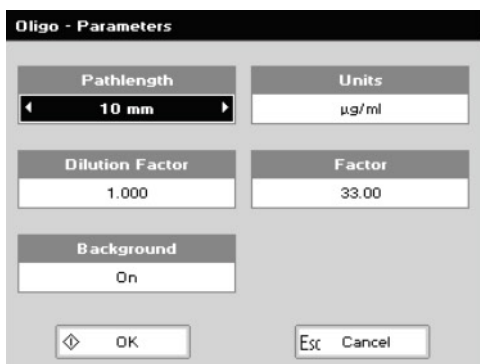
Results

The results are displayed on the screen.

Return to the Applications screen using the escape key OR display the options menu using the options key (see the Additional Options section).

Oligo

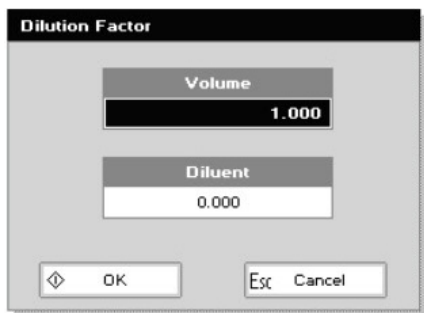
The Oligo application is accessed from the Nucleic Acids screen using the '3' key. It can be used to perform Oligonucleotide quantification measurements at 260 nm and to assess sample quality using A260/A280 and A260/A230 absorbance (A) ratios.



Step 1

Set the pathlength; "10 mm", "5 mm", "1 mm", "0.5 mm", "0.2 mm", or "0.125 mm", using the left and right arrow keys.

USER INTERFACE



Step 2

Press the down arrow, enter any dilution factor to be applied to the absorbance measurement of a value of up to four significant figures, using the alphanumeric keys.

OR

Open the dilution factor options using the options key:

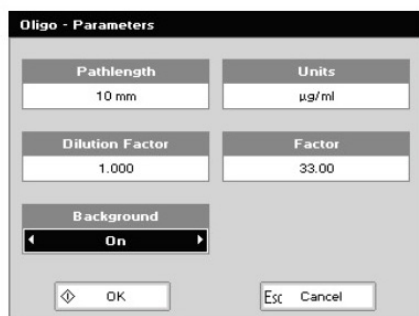
Set the initial sample volume of a value of up to four significant figures, using the alphanumeric keys.

Press the down arrow, set the amount of diluent added to the initial volume of a value of up to four significant figures, using the alphanumeric keys.

Implement the changes and return to the parameters screen by selecting "OK" using the confirm key.

OR

Reject the changes and return to the parameters screen by selecting "Cancel" using the escape key.



Step 3

Press the down arrow, set the background, "On" or "Off", using the left and right arrow keys.

Step 4

Press the down arrow, select one of the predefined units, "µg/ml", "ng/µl", "µg/µl", or "pmol/µl" using the left and right arrow keys.

For each predefined weight per volume unit a default factor is applied, "33.00" for "µg/ml" and "ng/µl", or "0.033" for "µg/µl".

OR

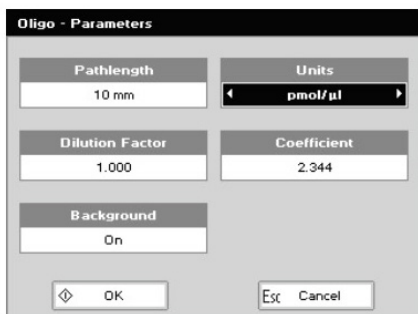
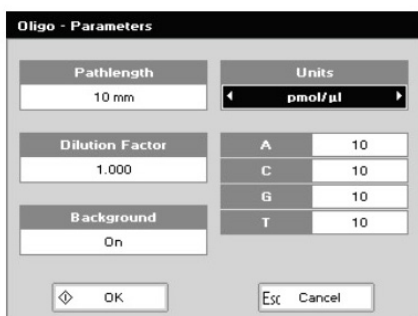
Press the down arrow and enter a custom value of up to four significant figures, using the alphanumeric keys.

For the "pmol/µl" unit two further options are available;

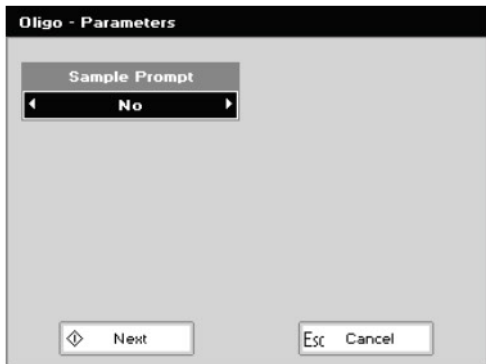
Either define the number of each base within the oligonucleotide sequence.

OR

Define the oligonucleotide's attenuation coefficient.



USER INTERFACE



Step 5

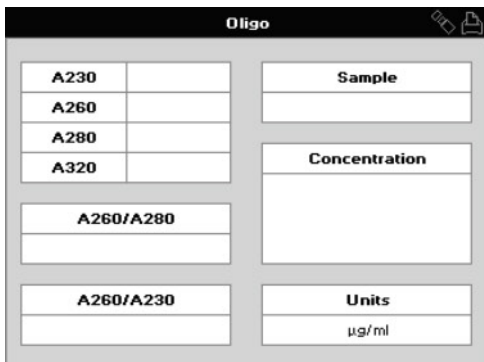
Proceed to the second parameter screen by selecting “OK” using the confirm key.

OR

Return to the Nucleic Acids screen by selecting “Cancel” using the escape key.

Step 6

Set the sample prompt, “Yes” or “No”, to bring up the sample number option before each measurement, using the left and right arrow keys.

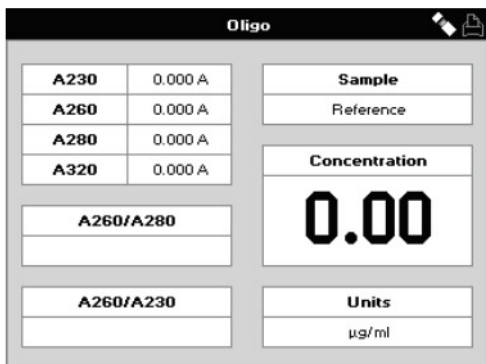


Step 7

Proceed to the measurement screen by selecting “OK” using the confirm key.

OR

Return to the previous parameter screen by selecting “Cancel” using the escape key.



Step 8

Insert the reference sample then take a reference measurement using the reference key.

The acquired reference sample baseline will be applied to all subsequent sample measurements.

USER INTERFACE

The screenshot shows a software interface titled "Fluorescent Dye - Parameters". It contains several input fields and buttons. The "Dye 1 Name" field is a dropdown menu currently showing "Cy2". The "Correction Factor" field contains the value "0.08". The "λ Max" field contains "489 nm". The "Extinction Coefficient" field contains "150.0 E+3". At the bottom left is a "Next" button with a right-pointing arrow, and at the bottom right is an "Est Cancel" button.

Step 9

Replace the previous sample with a test sample then take a sample measurement using the confirm key.

Repeat for all samples.

WARNING!

Previous measurements cannot be recalled so should be printed or output to a USB flash drive before any subsequent samples are measured.

Results

The results are displayed on the screen.

Return to the Applications screen using the escape key OR display the options menu using the options key (see the Additional Options section).

USER INTERFACE

Fluorescent Dye

The Fluorescent Dye application is accessed from the Nucleic Acids screen using the '4' key. It can be used to assess the fluorescent labelling efficiency of nucleic acid probes, based on the absorbance prior to their use in microarrays.

The screenshot shows the 'Fluorescent Dye - Parameters' screen. The 'Dye 1 Name' is set to 'Cy2' with left and right arrow keys. The 'Correction Factor' is 0.08. The 'λ Max' is 489 nm. The 'Extinction Coefficient' is 150.0 E+3. At the bottom, there are 'Next' and 'Esc Cancel' buttons.

Step 1

Select one of the predefined dyes using the left and right arrow keys.

Each dye has fixed associated parameters:

DYE 1 NAME	λ MAX	EXTINCTION COEFFICIENT	CORRECTION FACTOR
CY2	489 NM	150 E+3	0.08
CY3	550 NM	150 E+3	0.06
CY3B	558 NM	130 E+3	0.06
CY3.5	581 NM	150 E+3	0.14
CY5	647 NM	250 E+3	0.02
CY5.5	675 NM	190 E+3	0.15
CY7	747 NM	200 E+3	0.04
HYPER5	660 NM	110 E+3	0.25
FLUORESCCEIN	494 NM	92.3 E+3	0.32
ALEXA FLUOR 350	346 NM	19.0 E+3	0.25
ALEXA FLUOR 488	495 NM	71.0 E+3	0.30
ALEXA FLUOR 532	532 NM	81.0 E+3	0.24
ALEXA FLUOR 546	556 NM	104 E+3	0.21
ALEXA FLUOR 555	555 NM	150 E+3	0.04
ALEXA FLUOR 568	578 NM	91.3 E+3	0.45
ALEXA FLUOR 594	590 NM	73.0 E+3	0.43
ALEXA FLUOR 647	650 NM	239 E+3	0.00
ALEXA FLUOR 660	663 NM	110 E+3	0.00
ALEXA FLUOR 680	679 NM	184 E+3	0.00

The screenshot shows the 'Fluorescent Dye - Parameters' screen in 'Custom' mode. The 'Dye 1 Name' is 'Custom' with left and right arrow keys. The 'Correction Factor' is 0.00. The 'λ Max' is 679 nm. The 'Extinction Coefficient' is 184.0 E+3. There is a 'Custom Dye Name' field which is currently empty. At the bottom, there are 'Next' and 'Esc Cancel' buttons.

OR

Select the custom dye setting and define the "λ Max", 200 to 999 nm, Molar "Extinction Coefficient", 0.001 – 9999 E+3, A260 "Correction Factor", 0.01 to 9999, and a 16-digit "Custom Dye Name".

USER INTERFACE

Fluorescent Dye - Parameters

Dye 2 Name Cy3	Correction Factor 0.08
λ Max 550 nm	
Extinction Coefficient 150.0 E+3	

Next Esc Cancel

Step 2

Proceed to the second parameter screen by selecting “Next” using the confirm key.

OR

Return to the Nucleic Acids screen by selecting “Cancel” using the escape key.

Step 3

Select a second dye as detailed for the first dye in step 1.

OR

Select “none” to omit a second dye.

Fluorescent Dye - Parameters

Pathlength 10 mm	Dilution Factor 1.000
Background Correction Off	Volume (μl) 2.000
	Sample Prompt No

Next Esc Cancel

Step 4

Proceed to the third parameter screen by selecting “Next” using the confirm key.

OR

Return to the previous parameter screen by selecting “Cancel” using the escape key.

Step 5

Set the pathlength; “10 mm”, “5 mm”, “1 mm”, “0.5 mm”, “0.2 mm”, or “0.125 mm”, using the left and right arrow keys.

Step 6

Press the down arrow, set the background, “On” or “Off”, using the left and right arrow keys.

For background set to “On”, press the down arrow and set the background wavelength, 202 to 997, using the left and right arrow or alphanumeric keys.

Dilution Factor

Volume 1.000
Diluent 0.000

OK Esc Cancel

Step 7

Press the down arrow, enter any dilution factor to be applied to the absorbance measurement of a value of up to four significant figures, using the alphanumeric keys.

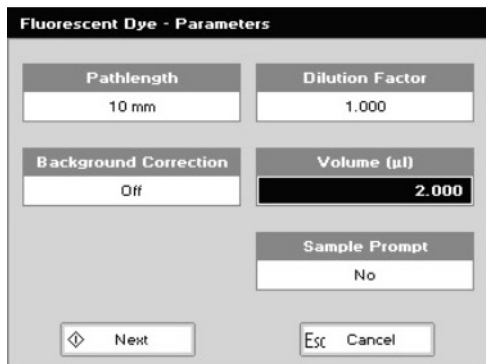
OR

Open the dilution factor options using the options key:

Set the initial sample volume of a value of up to four significant figures, using the alphanumeric keys.

Press the down arrow, set the amount of diluent added to the initial volume of a value of up to four significant figures, using the alphanumeric keys.

USER INTERFACE

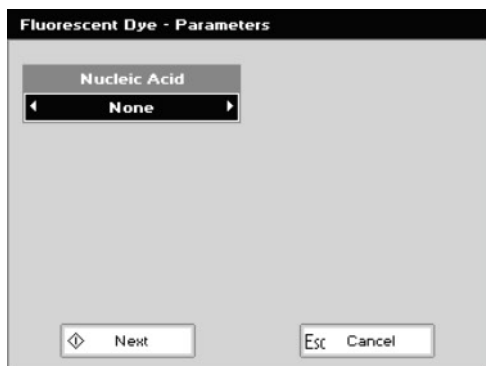


Step 8

Press the down arrow, set the volume a value of up to four significant figures, using the alphanumeric keys.

Step 9

Press the down arrow, set the sample prompt, “Yes” or “No”, to bring up the sample number option before each measurement, using the left and right arrow keys.



Step 10

Proceed to the fourth parameter screen by selecting “Next” using the confirm key.

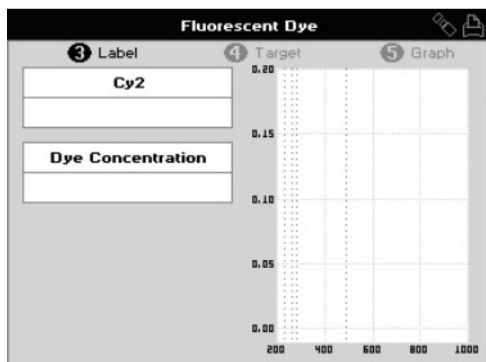
OR

Return to the previous parameter screen by selecting “Cancel” using the escape key.

Step 11

Set the nucleic acid target; “none”, “ssDNA (260 nm)”, “dsDNA (260 nm)”, “RNA (260 nm)”, “Cy Dye dUTP”, “Oligo (260 nm)”, or “Custom”, using the left and right arrow keys.

For the “Custom” target selection, press the down arrow and enter a factor of up to four significant figures, using the alphanumeric keys.



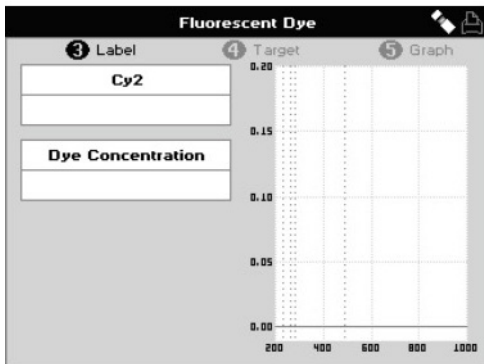
Step 12

Proceed to the measurement screen by selecting “Next” using the confirm key.

OR

Return to the previous parameter screen by selecting “Cancel” using the escape key.

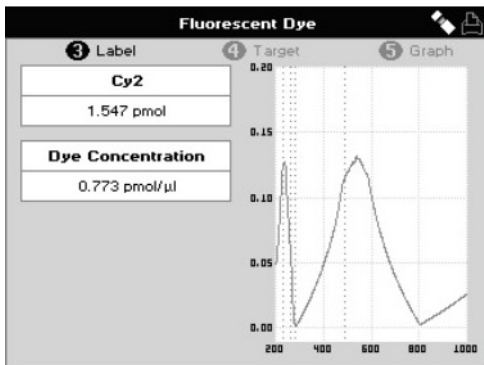
USER INTERFACE



Step 13

Insert the reference sample then take a reference measurement using the reference key.

The acquired reference sample baseline will be applied to all subsequent sample measurements.



Step 14

Replace the previous sample with a test sample then take a sample measurement using the confirm key.

Repeat for all samples.

WARNING!

Previous measurements cannot be recalled so should be printed or output to a USB flash drive before any subsequent samples are measured.

Results

The results are displayed on the screen.

Return to the Applications screen using the escape key OR display the options menu using the options key (see the Additional Options section).

USER INTERFACE

T_m Calculation

The melting temperature (T_m) Calculation application is accessed from the Nucleic Acids screen using the '5' key. It can be used to determine the theoretical melting point of a PCR primer at the measured concentration from its nucleotide base sequence, using the nearest-neighbour model.

The screenshot shows the 'Tm Calculation - Parameters' screen. It has a dark header with the title. Below the header are several input fields and buttons. The 'Base Type' field is a dropdown menu currently showing 'DNA'. The 'Buffer Molarity' field is a text input showing '0.100'. The 'Phosphorylated' field is a dropdown menu showing 'No'. The 'Counter Ion' field is a dropdown menu showing 'Na'. The 'Primer Conc.' field is a text input showing '1.000'. At the bottom, there are two buttons: 'Next' with a diamond icon and 'Esc Cancel'.

Step 1

Select the base type; “DNA” or “RNA”, using the left and right arrow keys.

Step 2

Press the down arrow, select if the nucleotides are phosphorylated; “No” or “Yes”, using the left and right arrow keys.

Step 3

Press the down arrow, enter the PCR primer concentration a value of up to four significant figures, using the alphanumeric keys.

Step 4

Press the down arrow, enter the buffer molarity (μM) a value of up to four significant figures, using the alphanumeric keys.

Step 5

Press the down arrow, select the counter ion present in the buffer; “Na”, “K”, “TEA”. “TEOA”, or “Other” up to a four digit value, using the left and right arrow keys.

For the “Other” counter ion selection, press the down arrow and enter the molecular weight (“Other MW”) up to four significant figures, using the alphanumeric keys.

The screenshot shows the 'Tm Calculation - Base Sequence' screen. It has a dark header with the title. Below the header are two input fields: 'Pathlength' showing '10 mm' and 'Sample Prompt' showing 'No'. Below these is a large text area labeled 'Base Sequence'. At the bottom, there are two buttons: 'OK' with a diamond icon and 'Esc Back'.

Step 6

Proceed to the base sequence screen by selecting “OK” using the confirm key.

OR

Return to the Applications screen by selecting “Cancel” using the escape key.

Step 7

Set the pathlength; “10 mm”, “5 mm”, “1 mm”, “0.5 mm”, “0.2 mm”, or “0.125 mm”, using the left and right arrow keys.

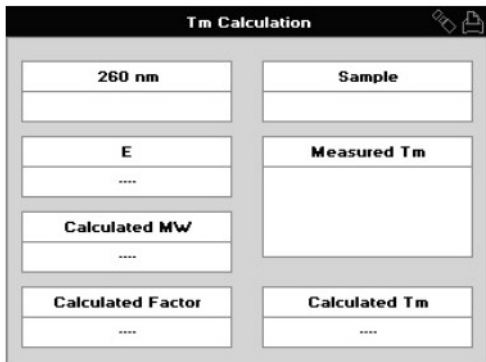
Step 8

Press the down arrow, enter the primer nucleotide base sequence of up to 60 bases using the ‘A’ (‘1’), ‘C’ (‘3’), ‘G’ (‘4’), and ‘T/U’ (‘6’) alphanumeric keys.

Step 9

Press the down arrow, set the sample prompt, “Yes” or “No”, to bring up the sample number option before each measurement, using the left and right arrow keys.

USER INTERFACE

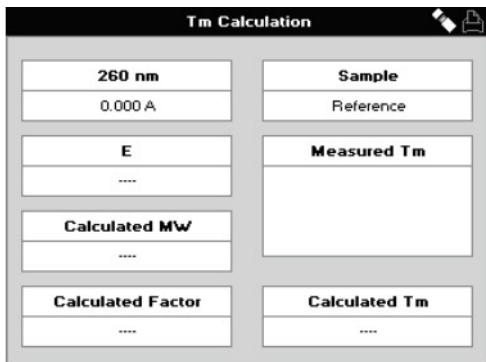


Step 10

Proceed to the measurement screen by selecting “OK” using the confirm key.

OR

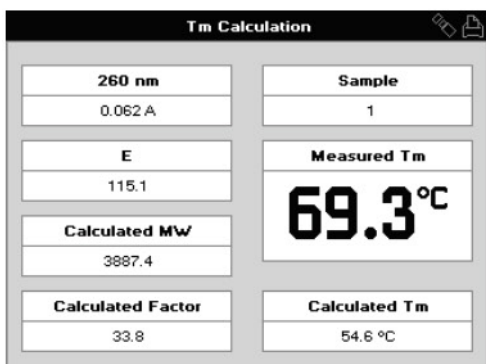
Return to the Applications screen by selecting “Back” using the escape key.



Step 11

Insert the reference sample then take a reference measurement using the reference key.

The acquired reference sample baseline will be applied to all subsequent sample measurements.



Step 12

Replace the previous sample with a test sample then take a sample measurement using the confirm key.

Repeat for all samples.

WARNING!

Previous measurements cannot be recalled so should be printed or output to a USB flash drive before any subsequent samples are measured.

Results

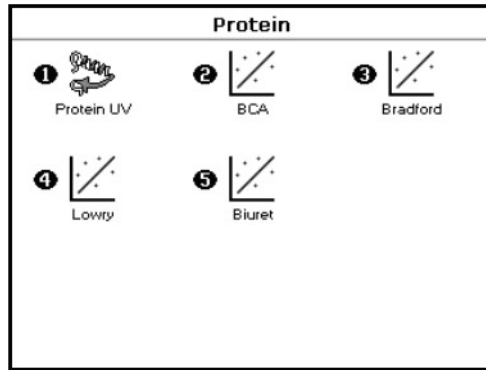
The results are displayed on the screen.

Return to the Applications screen using the escape key OR display the options menu using the options key (see the Additional Options section).

USER INTERFACE

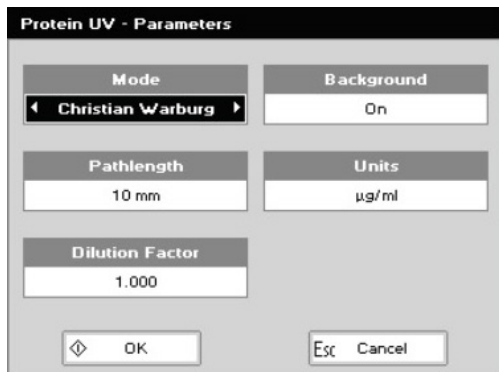
Protein

The Protein screen is accessed from the home screen using the '3' key. It contains predefined protein quantification methods. All calculation applied within the Protein applications are described in the Useful Calculation section.



Protein UV

The Protein UV application is accessed from the Protein screen using the '1' key. It can be used to perform Protein quantification measurements at 280 nm and to assess sample quality using A260/A280 and A260/A230 absorbance (A) ratios.

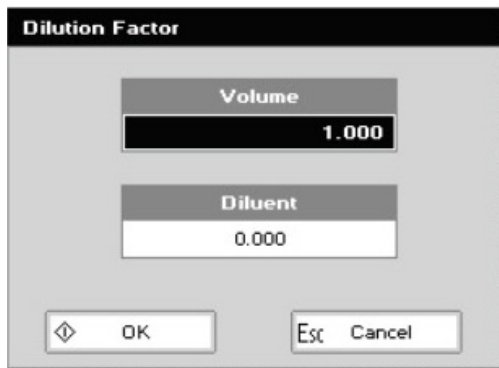


Step 1

Set the mode; "Christian Warburg", "BSA", "IgG", "Lysozyme", "Molar extinction", "Mass extinction", "E 1%", or "Custom", using the left and right arrow keys.

Step 2

Press the down arrow, set the pathlength; "10 mm", "5 mm", "1 mm", "0.5 mm", "0.2 mm", or "0.125 mm", using the left and right arrow keys.



Step 3

Press the down arrow, enter any dilution factor to be applied to the absorbance measurement of a value of up to four significant figures, using the alphanumeric keys.

OR

Open the dilution factor options using the options key: Set the initial sample volume of a value of up to four significant figures, using the alphanumeric keys. Press the down arrow, set the amount of diluent added

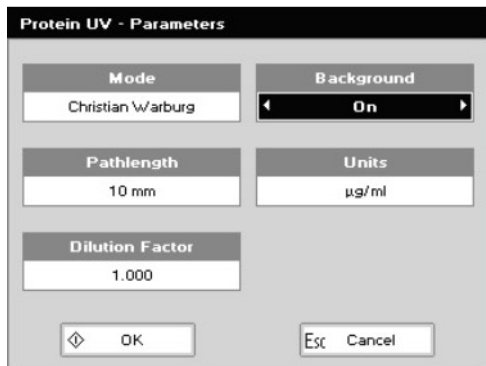
USER INTERFACE

to the initial volume of a value of up to four significant figures, using the alphanumeric keys.

Implement the changes and return to the parameters screen by selecting “OK” using the confirm key.

OR

Reject the changes and return to the parameters screen by selecting “Cancel” using the escape key.



Step 4

Press the down arrow, set the background, “On” or “Off”, using the left and right arrow keys.

Step 5

For “Christian Warburg”, “BSA”, “IgG”, and “Lysozyme” modes press the down arrow, select one of the predefined units, “µg/ml”, “ng/µl”, “µg/µl”, or “mg/ml”, using the left and right arrow keys.

OR

For the “Molar Extinction” mode press the down arrow, define the molar extinction coefficient (“AU l/mol ×1000”), then press the down arrow and define molecular weight (“MW kDa”) of the protein of interest.

OR

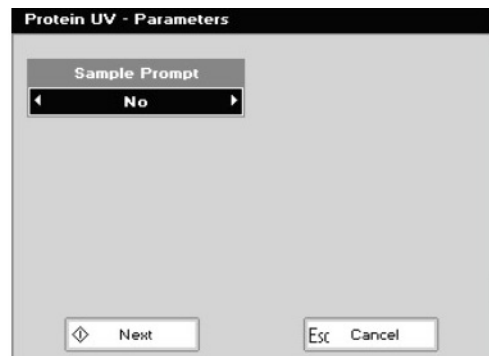
For the “Mass Extinction” mode press the down arrow, define the mass extinction coefficient (“AU l/g”) of the protein of interest.

OR

For the “E 1%” mode press the down arrow, define the 1% w/v extinction coefficient (“E 1%”) of the protein of interest.

OR

For the “Custom” mode press the down arrow, define the factors to apply to the absorbance measurements at 260 and 280 nm.



Step 6

Proceed to the second parameters screen by selecting “OK” using the confirm key.

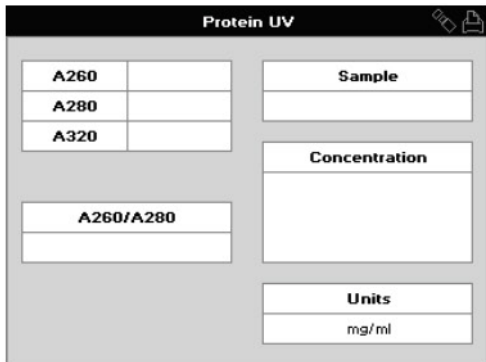
OR

Return to the Applications screen by selecting “Cancel” using the escape key.

Step 7

Set the sample prompt, “Yes” or “No”, to bring up the sample number option before each measurement, using the left and right arrow keys.

USER INTERFACE

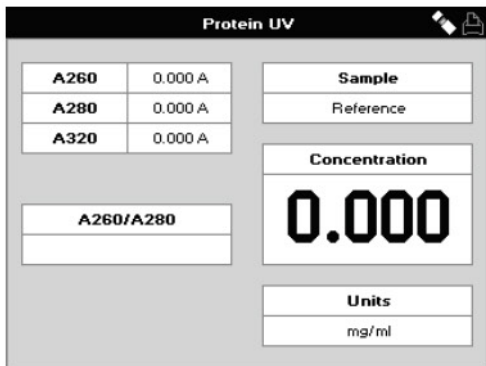


Step 8

Proceed to the measurement screen by selecting "OK" using the confirm key.

OR

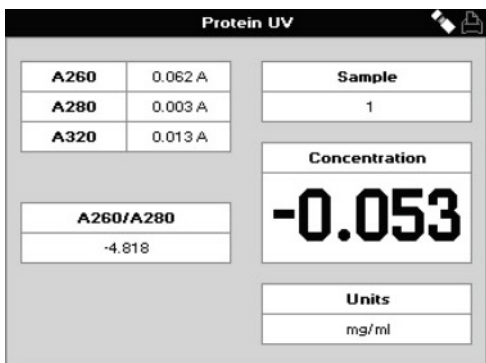
Return to the Applications screen by selecting "Cancel" using the escape key.



Step 9

Insert the reference sample then take a reference measurement using the reference key.

The acquired reference sample baseline will be applied to all subsequent sample measurements.



Step 10

Replace the previous sample with a test sample then take a sample measurement using the confirm key.

Repeat for all samples.

WARNING!

Previous measurements cannot be recalled so should be printed or output to a USB flash drive before any subsequent samples are measured.

Results

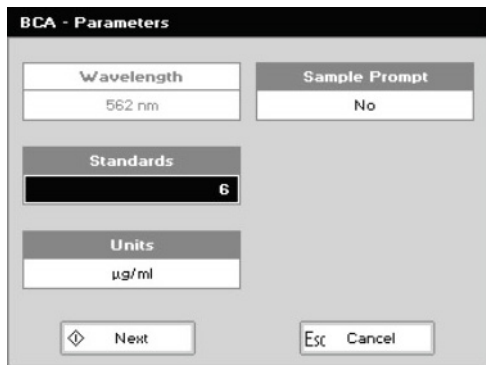
The results are displayed on the screen.

Return to the Applications screen using the escape key OR display the options menu using the options key (see the Additional Options section).

USER INTERFACE

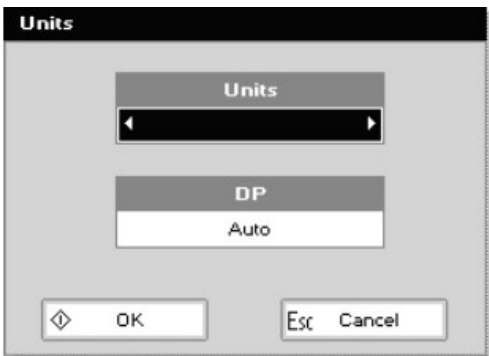
BCA

The BCA application is accessed from the Protein screen using the '2' key. It can be used to perform a BCA protein quantification assay based on the standard curve application.



Step 1

Set the number of standard samples of known concentration, 1 to 9, using the left and right arrow or alphanumeric keys.



Step 2

Press the down arrow, enter up to 8-digits to define the units that the primary result will be reported in.

OR

Open the units options using the options key:

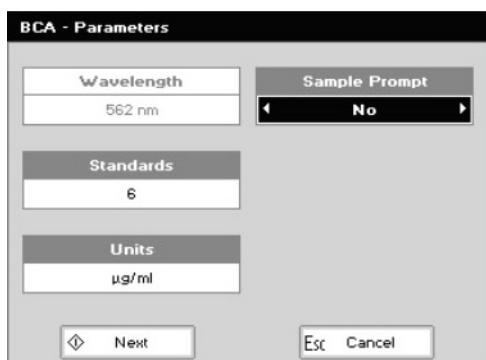
Select one of the predefined units; "µg/ml", "µg/µl", "pmol/µl", "mg/dl", "mmol/l", "µmol/l", "g/l", "mg/l", "µg/l", "U/l", "%", "ppm", "ppb", or "conc", using the left and right arrow keys.

Press the down arrow, select the number of decimal places to display the concentration in, "Auto", "0", "1", or "2", using the left and right arrow keys.

Implement the changes and return to the parameters screen by selecting "OK" using the confirm key.

OR

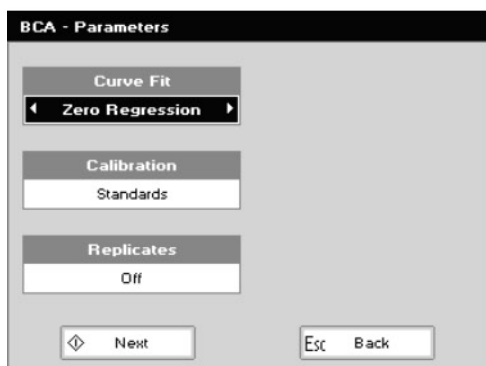
Reject the changes and return to the parameters screen by selecting "Cancel" using the escape key.



Step 3

Press the down arrow, set the sample prompt, "Yes" or "No", to bring up the sample number option before each measurement, using the left and right arrow keys.

USER INTERFACE



Step 4

Proceed to the second parameters screen by selecting “Next” using the confirm key.

OR

Return to the Applications screen by selecting “Cancel” using the escape key.

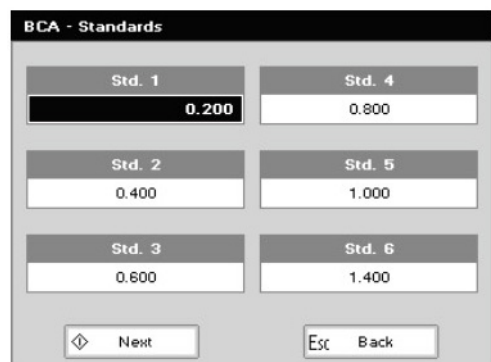
Step 5

Select the curve fit, “Zero Regression”, “2nd Order Polynomial”, “Regression”, “Interpolation”, or “Cubic Spline”, using the left and right arrow keys.

Step 6

Press the down arrow, select the source of the calibration, “Standards”, “Manual”, or “New Standards”, using the left and right arrow keys.

For “Standards” and “New Standards” calibration, press the down arrow and select the number of standard sample replicates, “Off”, “2”, or “3”, using the left and right arrow keys.



Step 7

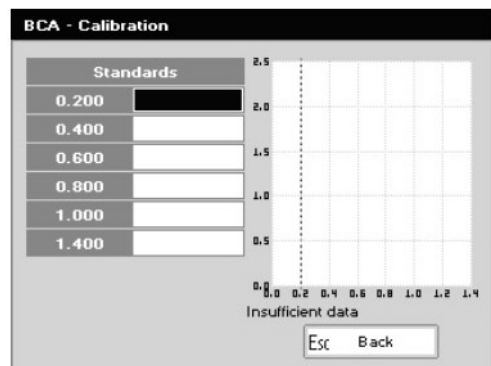
Proceed to the Standards screen by selecting “Next” using the confirm key.

OR

Return to the previous parameters screen by selecting “Back” using the escape key.

Step 8

Enter a value of up to four significant figures, for the known concentration of each standard to be used using the alphanumeric keys.



Step 9

Proceed to the Calibration screen by selecting “Next” using the confirm key.

OR

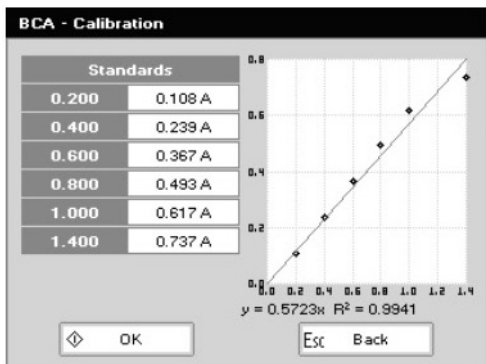
Return to the previous parameters screen by selecting “Back” using the escape key.

Step 10

Insert the reference sample then take a reference measurement using the reference key.

The acquired reference sample baseline will be applied to all subsequent standard and test sample

USER INTERFACE



Step 11

Replace the previous sample with the first standard sample then take a sample measurement using the confirm key.

Repeat for all standards samples.

BCA

Wavelength 562 nm	Sample Reference
Absorbance 0.000 A	Concentration
Curve Fit Zero Regression	Units ug/ml

Step 12

Proceed to the measurement screen by selecting "OK" using the confirm key.

OR

Return to the standards screen by selecting "Back" using the escape key.

BCA

Wavelength 562 nm	Sample 1
Absorbance 0.239 A	Concentration 0.418
Curve Fit Zero Regression	Units ug/ml

Step 13

Insert a test sample then take a sample measurement using the confirm key.

Repeat for all samples.

WARNING!

Previous measurements cannot be recalled so should be printed or output to a USB flash drive before any subsequent samples are measured.

Results

The results are displayed on the screen.

Return to the Applications screen using the escape key OR display the options menu using the options key (see the Additional Options section).

USER INTERFACE

Bradford

The Bradford application is accessed from the Protein screen using the '3' key. It can be used to perform a Bradford protein quantification assay based on the standard curve application.

Bradford - Parameters

Wavelength	595 nm	Sample Prompt	No
Standards	6		
Units	µg/ml		

Next Esc Cancel

Step 1

Set the number of standard samples of known concentration, 1 to 9, using the left and right arrow or alphanumeric keys.

Units

Units	
DP	Auto

OK Esc Cancel

Step 2

Press the down arrow, enter up to 8-digits to define the units that the primary result will be reported in.

OR

Open the units options using the options key:

Select one of the predefined units; “µg/ml”, “µg/µl”, “pmol/µl”, “mg/dl”, “mmol/l”, “µmol/l”, “g/l”, “mg/l”, “µg/l”, “U/l”, “%”, “ppm”, “ppb”, or “conc”, using the left and right arrow keys.

Press the down arrow, select the number of decimal places to display the concentration in, “Auto”, “0”, “1”, or “2”, using the left and right arrow keys.

Implement the changes and return to the parameters screen by selecting “OK” using the confirm key.

OR

Reject the changes and return to the parameters screen by selecting “Cancel” using the escape key.

Bradford - Parameters

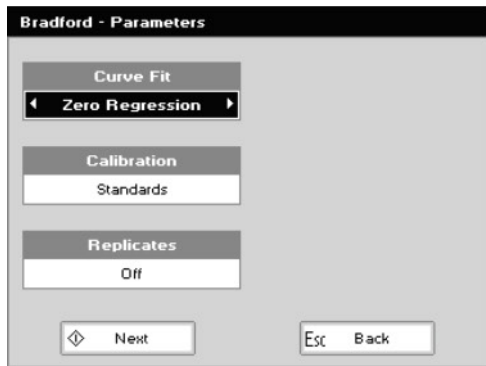
Wavelength	595 nm	Sample Prompt	No
Standards	6		
Units	µg/ml		

Next Esc Cancel

Step 3

Press the down arrow, set the sample prompt, “Yes” or “No”, to bring up the sample number option before each measurement, using the left and right arrow keys.

USER INTERFACE



Step 4

Proceed to the second parameters screen by selecting “Next” using the confirm key.

OR

Return to the Applications screen by selecting “Cancel” using the escape key.

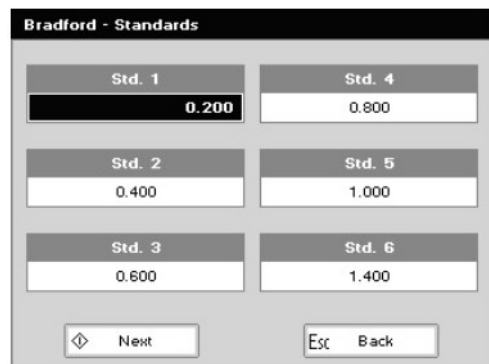
Step 5

Select the curve fit, “Zero Regression”, “2nd Order Polynomial”, “Regression”, “Interpolation”, or “Cubic Spline”, using the left and right arrow keys.

Step 6

Press the down arrow, select the source of the calibration, “Standards”, “Manual”, or “New Standards”, using the left and right arrow keys.

For “Standards” and “New Standards” calibration, press the down arrow and select the number of standard sample replicates, “Off”, “2”, or “3”, using the left and right arrow keys.



Step 7

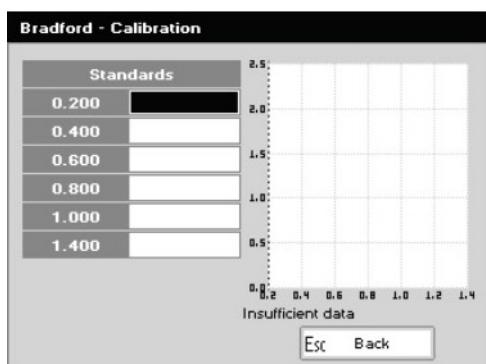
Proceed to the Standards screen by selecting “Next” using the confirm key.

OR

Return to the previous parameters screen by selecting “Back” using the escape key.

Step 8

Enter a value of up to four significant figures, for the known concentration of each standard to be used using the alphanumeric keys.



Step 9

Proceed to the Calibration screen by selecting “Next” using the confirm key.

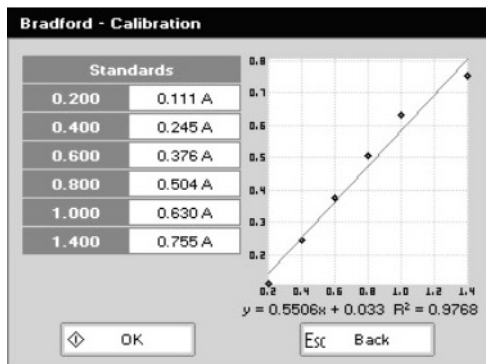
OR

Return to the previous parameters screen by selecting “Back” using the escape key.

Step 10

Insert the reference sample then take a reference measurement using the reference key.

USER INTERFACE



Step 11

Replace the previous sample with the first standard sample then take a sample measurement using the confirm key.

Repeat for all standards samples.

The measurement screen contains the following fields:

- Wavelength: 595 nm
- Sample: Reference
- Absorbance: 0.000 A
- Concentration: (empty)
- Curve Fit: Regression
- Units: $\mu\text{g/ml}$

Step 12

Proceed to the measurement screen by selecting "OK" using the confirm key.

OR

Return to the standards screen by selecting "Back" using the escape key.

The measurement screen shows the final result:

- Wavelength: 595 nm
- Sample: 1
- Absorbance: 0.245 A
- Concentration: **0.385**
- Curve Fit: Regression
- Units: $\mu\text{g/ml}$

Step 13

Insert a test sample then take a sample measurement using the confirm key.

Repeat for all samples.

WARNING!

Previous measurements cannot be recalled so should be printed or output to a USB flash drive before any subsequent samples are measured.

Results

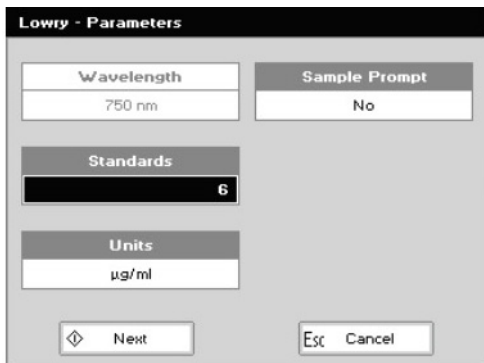
The results are displayed on the screen.

Return to the Applications screen using the escape key OR display the options menu using the options key (see the Additional Options section).

USER INTERFACE

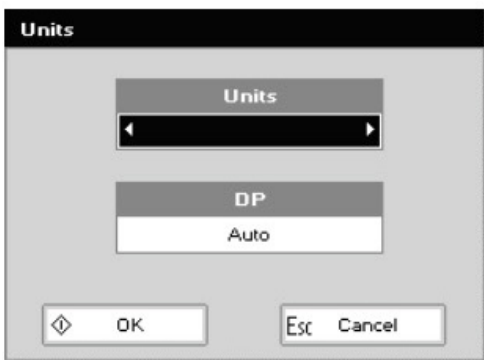
Lowry

The Lowry application is accessed from the Protein screen using the '4' key. It can be used to perform a Lowry protein quantification assay based on the standard curve application.



Step 1

Set the number of standard samples of known concentration, 1 to 9, using the left and right arrow or alphanumeric keys.



Step 2

Press the down arrow, enter up to 8-digits to define the units that the primary result will be reported in.

OR

Open the units options using the options key:

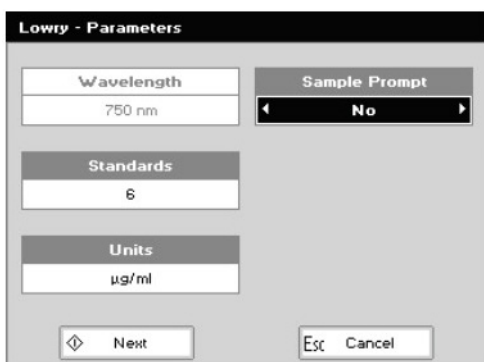
Select one of the predefined units; “µg/ml”, “µg/µl”, “pmol/µl”, “mg/dl”, “mmol/l”, “µmol/l”, “g/l”, “mg/l”, “µg/l”, “U/l”, “%”, “ppm”, “ppb”, or “conc”, using the left and right arrow keys.

Press the down arrow, select the number of decimal places to display the concentration in, “Auto”, “0”, “1”, or “2”, using the left and right arrow keys.

Implement the changes and return to the parameters screen by selecting “OK” using the confirm key.

OR

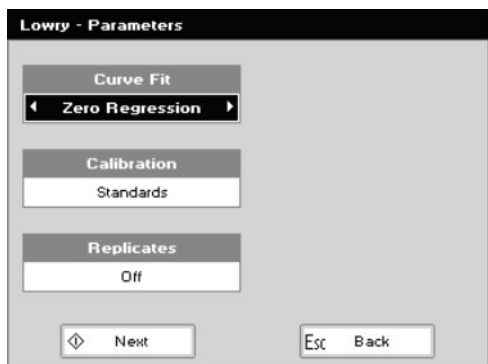
Reject the changes and return to the parameters screen by selecting “Cancel” using the escape key.



Step 3

Press the down arrow, set the sample prompt, “Yes” or “No”, to bring up the sample number option before each measurement, using the left and right arrow keys.

USER INTERFACE



Step 4

Proceed to the second parameters screen by selecting "Next" using the confirm key.

OR

Return to the Applications screen by selecting "Cancel" using the escape key.

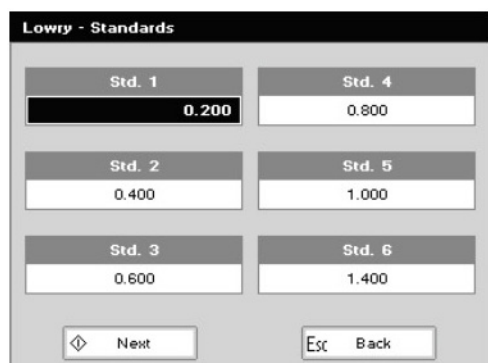
Step 5

Select the curve fit, "Zero Regression", "2nd Order Polynomial", "Regression", "Interpolation", or "Cubic Spline", using the left and right arrow keys.

Step 6

Press the down arrow, select the source of the calibration, "Standards", "Manual", or "New Standards", using the left and right arrow keys.

For "Standards" and "New Standards" calibration, press the down arrow and select the number of standard sample replicates, "Off", "2", or "3", using the left and right arrow keys.



Step 7

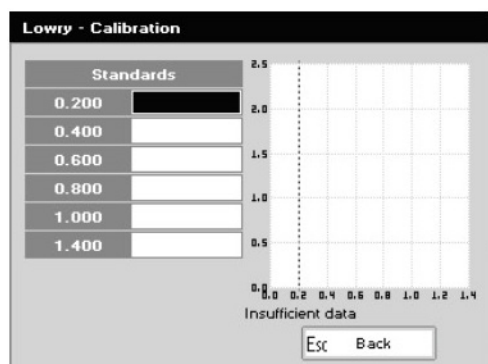
Proceed to the Standards screen by selecting "Next" using the confirm key.

OR

Return to the previous parameters screen by selecting "Back" using the escape key.

Step 8

Enter a value of up to four significant figures, for the known concentration of each standard to be used using the alphanumeric keys.



Step 9

Proceed to the Calibration screen by selecting "Next" using the confirm key.

OR

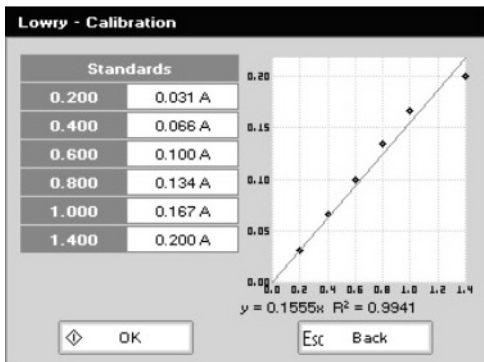
Return to the previous parameters screen by selecting "Back" using the escape key.

Step 10

Insert the reference sample then take a reference measurement using the reference key.

The acquired reference sample baseline will be applied to all subsequent standard and test sample measurements.

USER INTERFACE



Step 11

Replace the previous sample with the first standard sample then take a sample measurement using the confirm key.

Repeat for all standards samples.

The measurement screen shows the following settings:

- Wavelength: 750 nm
- Sample: Reference
- Absorbance: 0.000 A
- Concentration: (empty field)
- Curve Fit: Zero Regression
- Units: $\mu\text{g/ml}$

Step 12

Proceed to the measurement screen by selecting "OK" using the confirm key.

OR

Return to the standards screen by selecting "Back" using the escape key.

The measurement screen shows the final result:

- Wavelength: 750 nm
- Sample: 1
- Absorbance: 0.066 A
- Concentration: **0.424**
- Curve Fit: Zero Regression
- Units: $\mu\text{g/ml}$

Step 13

Insert a test sample then take a sample measurement using the confirm key.

Repeat for all samples.

WARNING!

Previous measurements cannot be recalled so should be printed or output to a USB flash drive before any subsequent samples are measured.

Results

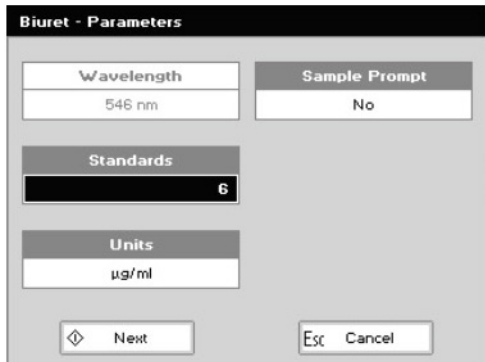
The results are displayed on the screen.

Return to the Applications screen using the escape key OR display the options menu using the options key (see the Additional Options section).

USER INTERFACE

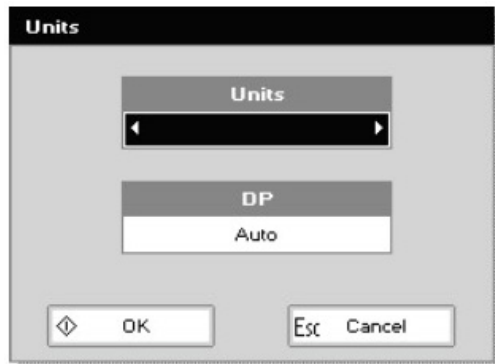
Biuret

The Biuret application is accessed from the Protein screen using the '5' key. It can be used to perform a Biuret protein quantification assay based on the standard curve application.



Step 1

Set the number of standard samples of known concentration, 1 to 9, using the left and right arrow or alphanumeric keys.



Step 2

Press the down arrow, enter up to 8-digits to define the units that the primary result will be reported in.

OR

Open the units options using the options key:

Select one of the predefined units; "µg/ml", "µg/µl", "pmol/µl", "mg/dl", "mmol/l", "µmol/l", "g/l", "mg/l", "µg/l", "U/l", "%", "ppm", "ppb", or "conc", using the left and right arrow keys.

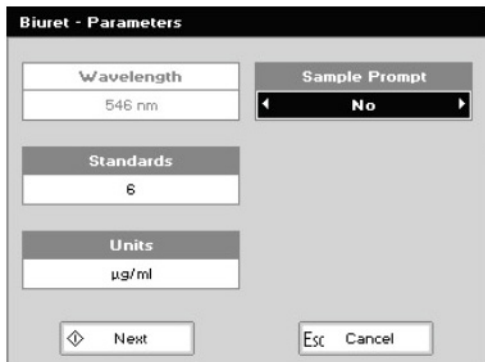
Press the down arrow, select the number of decimal places to display the concentration in, "Auto", "0", "1", or "2", using the left and right arrow keys.

Implement the changes and return to the parameters screen by selecting "OK" using the confirm key.

OR

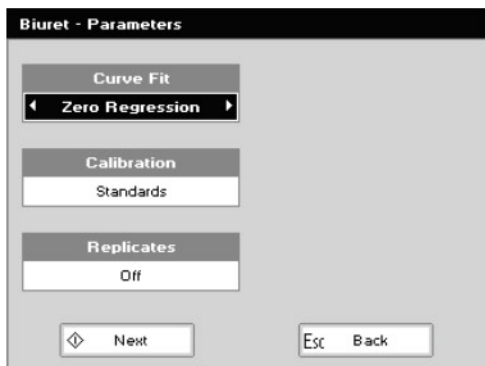
Reject the changes and return to the parameters screen by selecting "Cancel" using the escape key.

USER INTERFACE



Step 3

Press the down arrow, set the sample prompt, “Yes” or “No”, to bring up the sample number option before each measurement, using the left and right arrow keys.



Step 4

Proceed to the second parameters screen by selecting “Next” using the confirm key.

OR

Return to the Applications screen by selecting “Cancel” using the escape key.

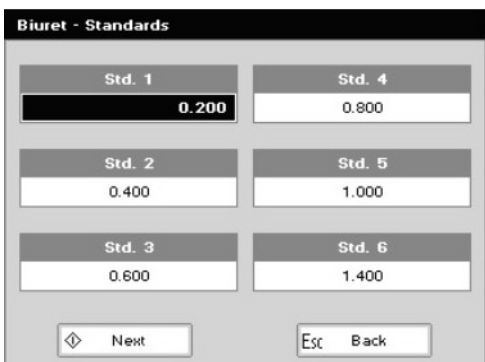
Step 5

Select the curve fit, “Zero Regression”, “2nd Order Polynomial”, “Regression”, “Interpolation”, or “Cubic Spline”, using the left and right arrow keys.

Step 6

Press the down arrow, select the source of the calibration, “Standards”, “Manual”, or “New Standards”, using the left and right arrow keys.

For “Standards” and “New Standards” calibration, press the down arrow and select the number of standard sample replicates, “Off”, “2”, or “3”, using the left and right arrow keys.



Step 7

Proceed to the Standards screen by selecting “Next” using the confirm key.

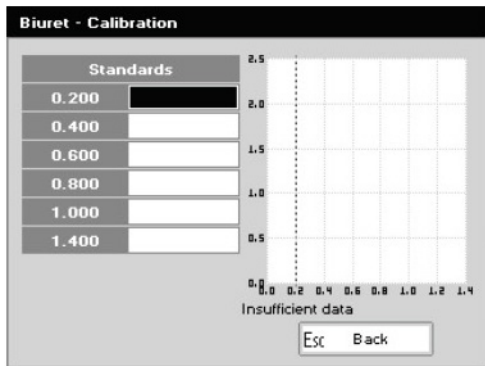
OR

Return to the previous parameters screen by selecting “Back” using the escape key.

Step 8

Enter a value of up to four significant figures, for the known concentration of each standard to be used using the alphanumeric keys.

USER INTERFACE



Step 9

Proceed to the Calibration screen by selecting “Next” using the confirm key.

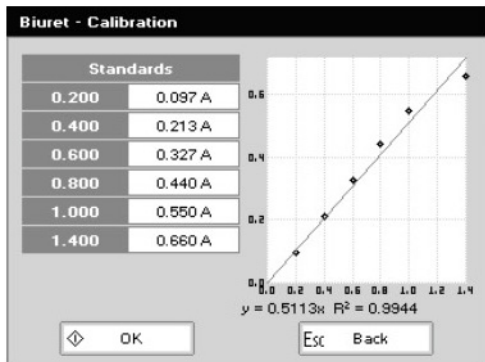
OR

Return to the previous parameters screen by selecting “Back” using the escape key.

Step 10

Insert the reference sample then take a reference measurement using the reference key.

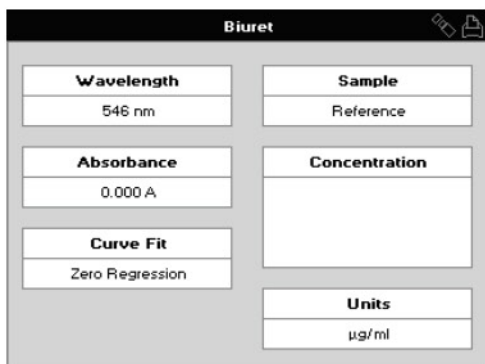
The acquired reference sample baseline will be applied to all subsequent standard and test sample measurements.



Step 11

Replace the previous sample with the first standard sample then take a sample measurement using the confirm key.

Repeat for all standards samples.



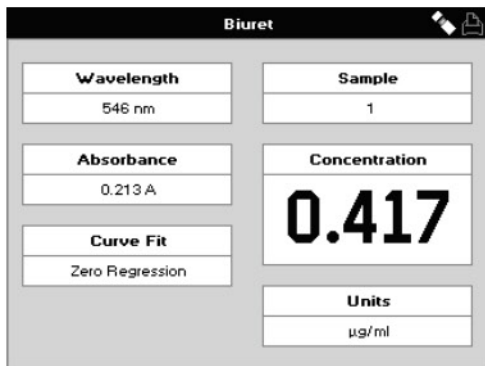
Step 12

Proceed to the measurement screen by selecting “OK” using the confirm key.

OR

Return to the standards screen by selecting “Back” using the escape key.

USER INTERFACE



Step 13

Insert a test sample then take a sample measurement using the confirm key.

Repeat for all samples.

WARNING!

Previous measurements cannot be recalled so should be printed or output to a USB flash drive before any subsequent samples are measured.

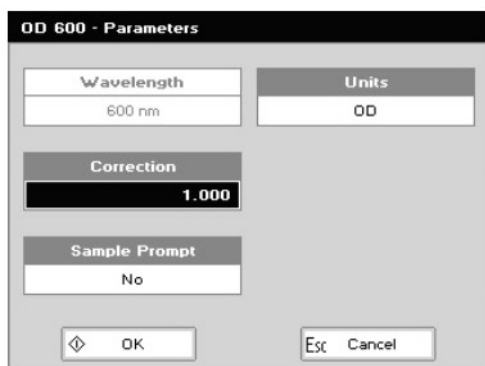
Results

The results are displayed on the screen.

Return to the Applications screen using the escape key OR display the options menu using the options key (see the Additional Options section).

OD 600

The OD 600 application is accessed from the home screen using the '4' key. It can be used to perform simple optical density measurements of microbial growth cultures.



Step 1

Enter any correction factor to be applied to the absorbance measurement of a value of up to four significant figures, using the alphanumeric keys.

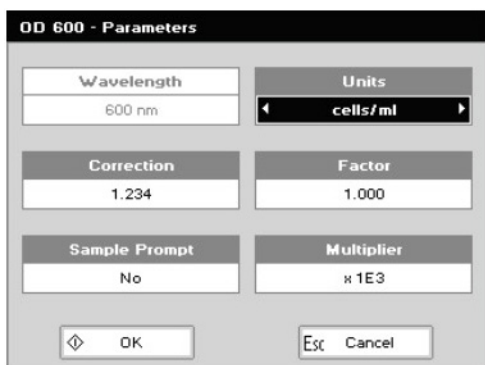
Step 2

Press the down arrow, set the sample prompt, "Yes" or "No", to bring up the sample number option before each measurement, using the left and right arrow keys.

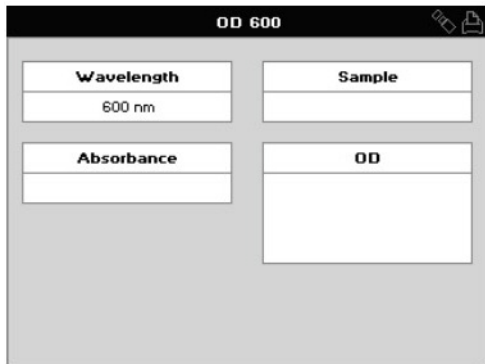
Step 3

Press the down arrow, set the units, "OD" or "cells/ml", using the left and right arrow keys.

For "cells/ml" units, press the down arrow and define the multiplication factor of a value of up to four significant figures, using the alphanumeric keys, then press the down arrow and set the multiplier, "x1E3" or "x1E6", using the left and right arrow keys.



USER INTERFACE

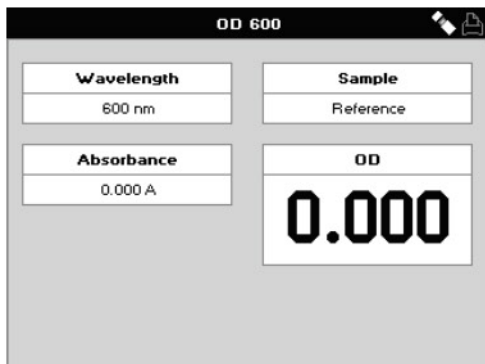


Step 4

Proceed to the measurement screen by selecting “OK” using the confirm key.

OR

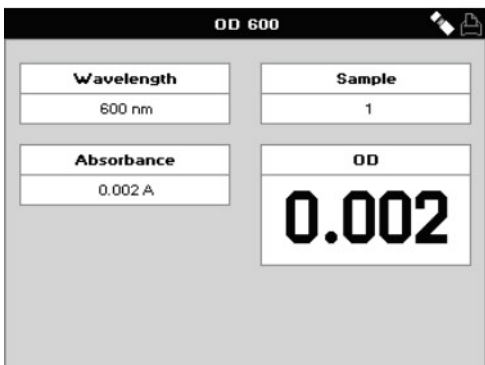
Return to the Applications screen by selecting “Cancel” using the escape key.



Step 5

Insert the reference sample then take a reference measurement using the reference key.

The acquired reference sample baseline will be applied to all subsequent sample measurements.



Step 6

Replace the previous sample with a test sample then take a sample measurement using the confirm key.

Repeat for all samples.

WARNING!

Previous measurements cannot be recalled so should be printed or output to a USB flash drive before any subsequent samples are measured.

Results

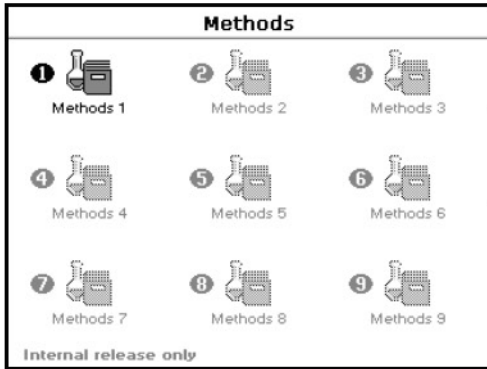
The results are displayed on the screen.

Return to the Applications screen using the escape key OR display the options menu using the options key (see the Additional Options section).

USER INTERFACE

Favourites, Methods, and USB Memory Stick

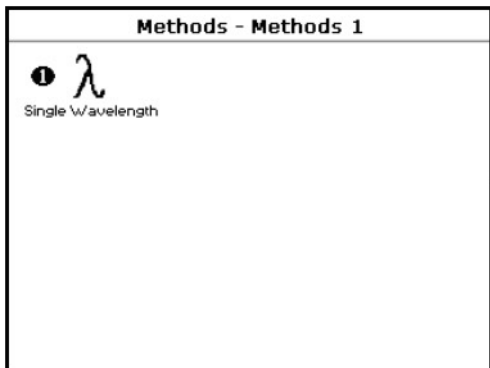
The Favourites, Methods, and USB Memory Stick screens are accessed from the home screen using the '5', '6', or '8' key respectively. They are directories to save custom methods to, using the options menu from the results screen (see the "Additional Options" section).



Select the methods subdirectory, where the custom method is saved, using the appropriate alphanumeric key.

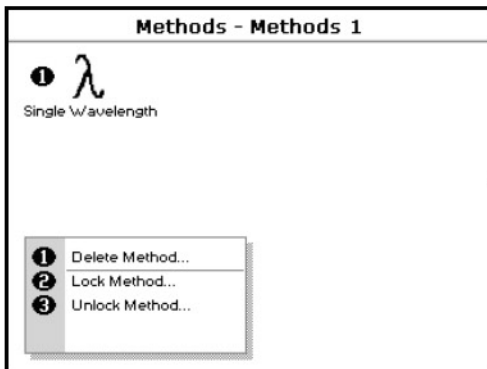
PLEASE NOTE

This only applies to the Methods screen. When selected from the home screen, the Favourites and Memory Stick screens directly display the saved custom method applications



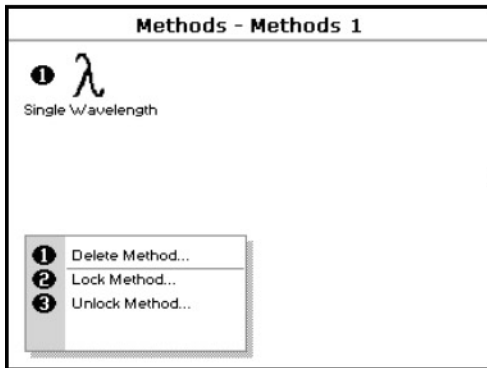
Once the directory containing the saved custom method application is accessed, there are several options available OR return to the home screen using the escape key.

1. To open a saved custom method application, use the appropriate alphanumeric key.

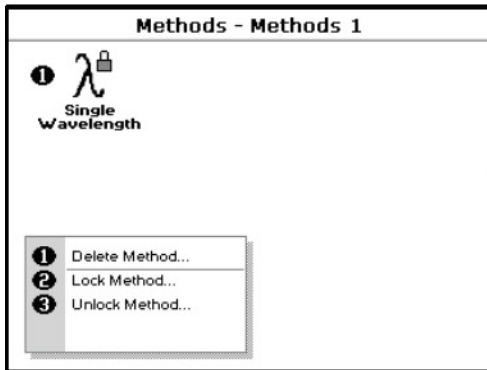


2. To delete a saved custom method application, use the options key then press the '1' key.

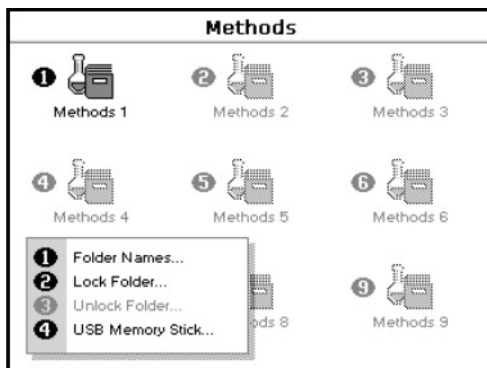
USER INTERFACE



3. To lock protect a saved custom method application, use the options key then press the '2' key.



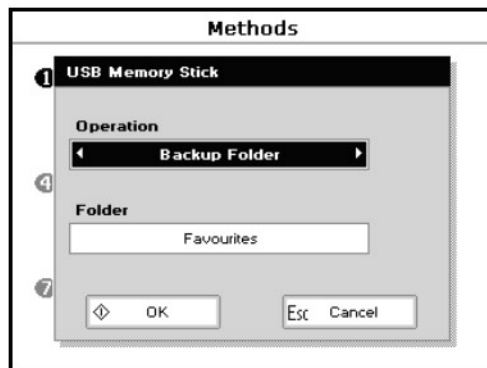
4. To unlock a protected saved custom method application, use the options key then press the '3' key.



5. To back up or restore custom method applications, insert a USB flash drive then use the options key then press the '4' key.

Select the operation, "Backup Folder", "Restore Folder", "Backup All Folders", or "Restore All Folders", using the left and right arrow keys.

For "Backup Folder" and "Restore Folder" operations, press the down arrow and select the folder to perform the operation on, "Favourites", "Methods 1", "Methods 2", "Methods 3", "Methods 4", "Methods 5", "Methods 6", "Methods 7", "Methods 8", or "Methods 9", using the left and right arrow keys.



Perform the operation and return to the Methods screen by selecting "OK" using the confirm key.

OR

Cancel the operation and return to the Methods screen by selecting "Cancel" using the escape key.

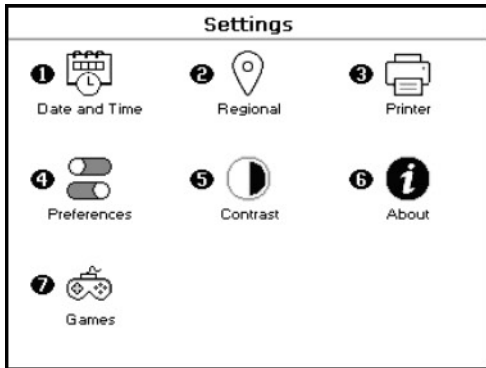
WARNING!

This operation cannot be undone and even locked methods are effected. Therefore, it is recommended to make a separate backup before restoring from a previous one.

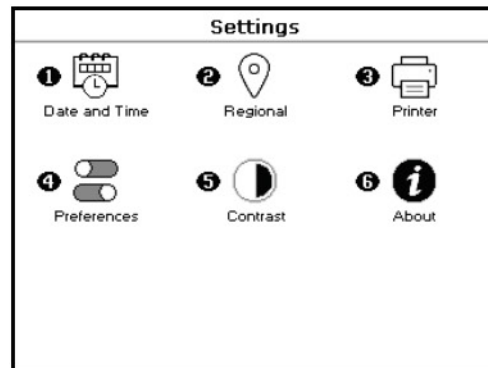
USER INTERFACE

Settings

The Settings screen is accessed from the home screen using the '7' key. It can be used to access application to adjust the instrument settings, such as date, time, language, and number format.



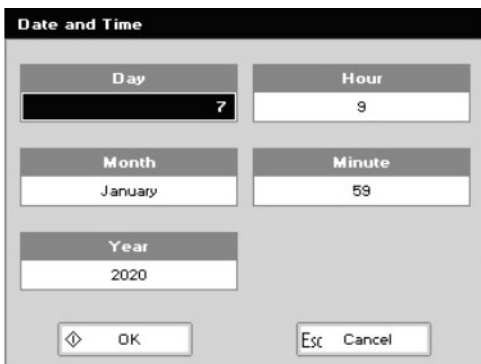
Default Settings screen displaying the Games application



Settings screen with the Games application hidden

Date and Time

The Date and Time application is accessed from the Settings screen using the '1' key. It can be used to adjust the date and time stamp applied to measurement data outputs.



Step 1

Set the day using the left and right arrow or alphanumeric keys.

Step 2

Press the down arrow, set the month using the left and right arrow keys.

Step 3

Press the down arrow, set the year using the left and right arrow or alphanumeric keys.

Step 4

Press the down arrow, set the hour (24-hour format) using the left and right arrow or alphanumeric keys.

Step 5

Press the down arrow, set the minute using the left and right arrow or alphanumeric keys.

Implement the changes and return to the Settings screen by selecting OK using the confirm key.

OR

Reject the changes and return to the Settings screen by selecting "Cancel" using the escape key.

USER INTERFACE

Regional

The Regional application is accessed from the Settings screen using the '2' key. It can be used to change the language and decimal separator number format.



Step 1

Set the language using the left and right arrow keys.

Step 2

Press the down arrow, set the number format decimal separator using the left and right arrow keys.

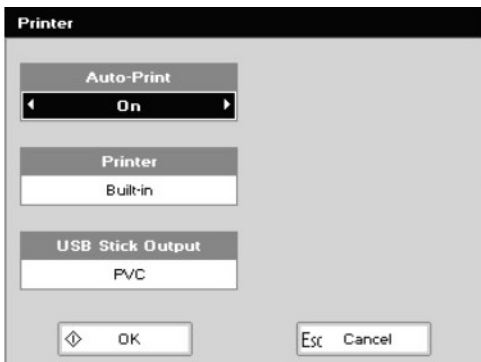
Implement the changes and return to the Settings screen by selecting OK using the confirm key.

OR

Reject the changes and return to the Settings screen by selecting "Cancel" using the escape key.

Printer

The Printer application is accessed from the Settings screen using the '3' key. It can be used to define the default printer and data output settings.



Step 1

Set auto-print "On" of "Off" using the left and right arrow keys.

Step 2

Press the down arrow, set the printer output to a "Built-in" printer, a computer via a USB cable ("Computer (USB)"), or a computer via Bluetooth ("Computer (Bluetooth)") depending on what is connected to the instrument, using the left and right arrow keys.

Step 3

Press the down arrow, set the USB file type for output to a connect USB flash drive ("PVC" or "TSV") using the left and right arrow keys.

Implement the changes and return to the Settings screen by selecting OK using the confirm key.

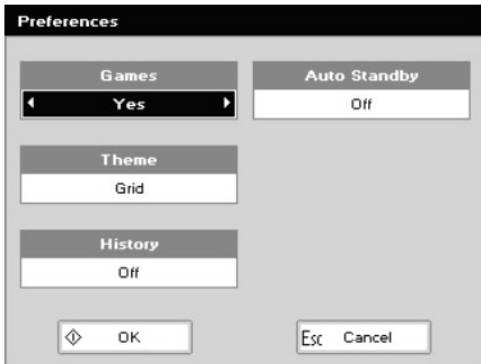
OR

Reject the changes and return to the Settings screen by selecting "Cancel" using the escape key.

USER INTERFACE

Preferences

The Preference application is accessed from the Settings screen using the '4' key. It can be used to set the access to the games application, the menu layout theme, parameter history, and standby settings.



Step 1

Set access to the games application or not, using the left and right arrow keys.

Step 2

Press the down arrow, set the application menus layout theme using the left and right arrow keys.

Step 3

Press the down arrow, set the parameter history to store application settings for future use or not, using the left and right arrow keys.

Step 4

Press the down arrow, set the auto standby interval using the left and right arrow keys.

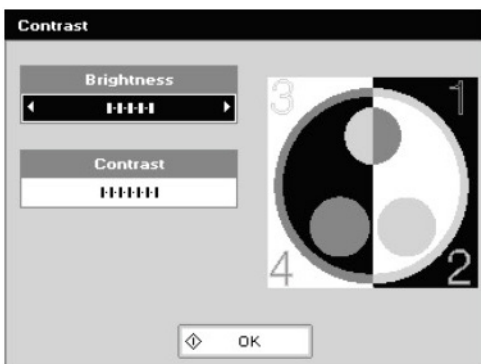
Implement the changes and return to the Settings screen by selecting OK using the confirm key.

OR

Reject the changes and return to the Settings screen by selecting "Cancel" using the escape key.

Contrast

The Contrast application is accessed from the Settings screen using the '5' key. It can be used to set the display brightness and contrast.



Step 1

Set the brightness using the left and right arrow keys.

Step 2

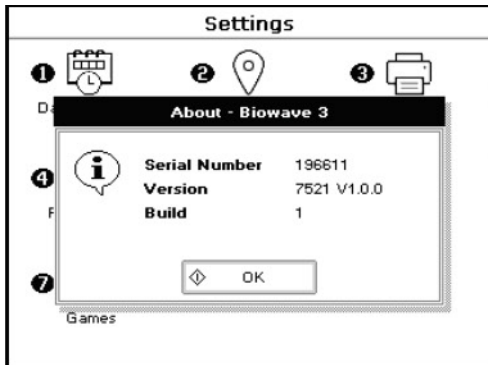
Press the down arrow, set the contrast using the left and right arrow keys.

Implement the changes and return to the Settings screen by selecting OK using the confirm key.

USER INTERFACE

About

The About pop-up window is opened from the Settings screen using the '6' key. The pop-up window displays information about the instrument, including its serial number, firmware version, and build.



Step 1

Set the brightness using the left and right arrow keys.

Step 2

Press the down arrow, set the contrast using the left and right arrow keys.

Implement the changes and return to the Settings screen by selecting OK using the confirm key.

Additional Options

Additional options are available from the measurement screen using the options key. The available options for key press 3, 4, 5, and 6 varies between applications.

Single Wavelength	Multi Wavelength	Concentration
<ul style="list-style-type: none"> 1 Parameters... 2 Print 3 Abs/%T 4 Print Graph 7 Sample Number... 8 Save Method... 9 Auto-Print ✓ 	<ul style="list-style-type: none"> 1 Parameters... 2 Print 4 Print Graph 7 Sample Number... 8 Save Method... 9 Auto-Print ✓ 	<ul style="list-style-type: none"> 1 Parameters... 2 Print 3 Graph 4 Run Standard... 7 Sample Number... 8 Save Method... 9 Auto-Print ✓
Standard Curve Absorbance Ratio	Wavescan	Kinetics
<ul style="list-style-type: none"> 1 Parameters... 2 Print 3 Graph 7 Sample Number... 8 Save Method... 9 Auto-Print ✓ 	<ul style="list-style-type: none"> 1 Parameters... 2 Print 3 Abs/%T 4 Peak Detection... 5 Add Peak 6 Graph Scale... 7 Sample Number... 8 Save Method... 9 Auto-Print ✓ 	<ul style="list-style-type: none"> 1 Parameters... 2 Print 3 Print Data 4 Set t0 At Cursor 5 Set t1 At Cursor 6 Slope 7 Sample Number... 8 Save Method... 9 Auto-Print ✓

Key Press

Description

1

Parameters – Return to method parameters screen.

2

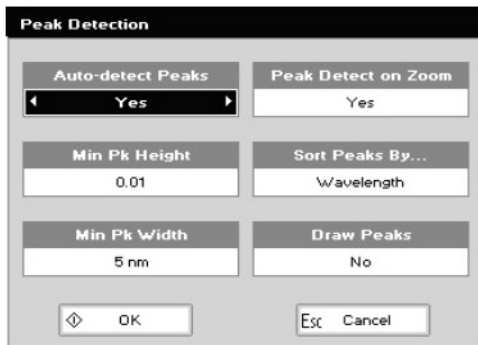
Print – Print result via selected method.

3

Abs/%T – Toggle between Absorbance and %T mode.
Graph – Displays the graph showing the measurements' local spectrum.
Print Data – Prints tabulated time-point data.

USER INTERFACE

4



Print Graph – Print graph via selected setting.

Run Standard – Re-run and standard sample measurement.

Peak Detection – Open the peak detection parameters to define the auto-peak detection settings.

Step 1

Set the method to auto-detect peaks or not using the left and right arrow keys.

For “Yes” to auto-detect peaks, press the down arrow and set the minimum peak height (0.00 to 9999) using the alphanumeric keys. Then press the down arrow and set the minimum peak width (1 to 190 nm) using the alphanumeric keys. Then press the down arrow and set to peak detect on zoom or not using the left and right arrow keys.

Step 2

Press the down arrow, set the sort by peaks by “Wavelength”, “Peak Height”, or “Peak Width”, using the left and right arrow keys.

Step 3

Press the down arrow, set to draw peaks or not using the left and right arrow keys.

Implement the changes and return to the parameters screen by selecting “OK” using the confirm key.

OR

Reject the changes and return to the parameters screen by selecting “Cancel” using the escape key.

Set t0 At Cursor – Sets the zero time-point of the slope at the current cursor position.

Target – Toggles to the “Target” measurement screen.

5

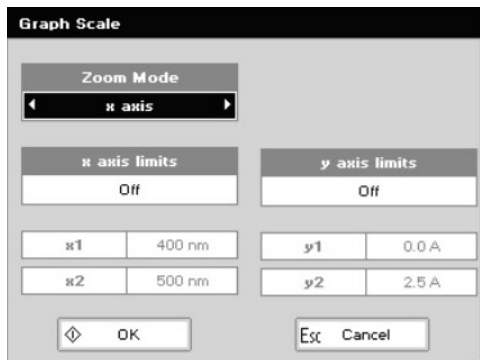
Add Peak – Adds the measurement at the current cursor position to the peak table.

Set tn At Cursor – Sets the final time-point of the slope at the current cursor position.

Graph – Toggles to the “Graph” screen showing the measurements’ local spectrum.

USER INTERFACE

6



Graph Scale – Open the graph scale parameters

Step 1

Set the zoom mode using the left and right arrow keys.

Step 2

Press the down arrow, apply x axis limits or not using the left and right arrow keys.

For “On” to x axis limits, press the down key and set the x-axis start value (“x1”) using the alphanumeric keys, then press the down arrow and set the x-axis end value (“x2”).

Step 3

Press the down arrow, apply y axis limits or not using the left and right arrow keys.

For “On” to y axis limits, press the down key and set the y-axis start value (“y1”) using the alphanumeric keys, then press the down arrow and set the y-axis end value (“y2”).

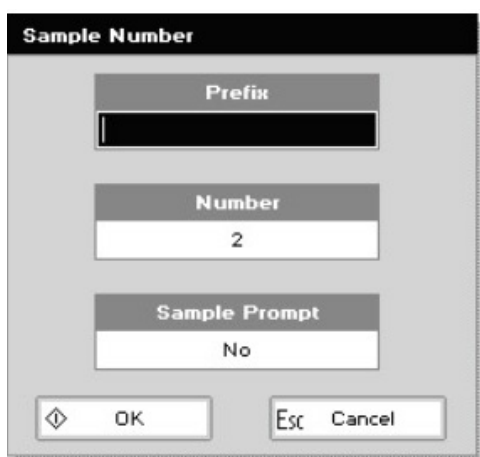
Implement the changes and return to the parameters screen by selecting “OK” using the confirm key.

OR

Reject the changes and return to the parameters screen by selecting “Cancel” using the escape key.

Slope – Fits a trend line across the time-course data or between the set t0 and tn points.

7



Sample Number...

Open sample number options.

Insert up to an 8-digit prefix using the alphanumeric keys.

Press the down arrow, insert a number up to 9999.

Press the down arrow, change the sample prompt setting using the left and right arrow keys.

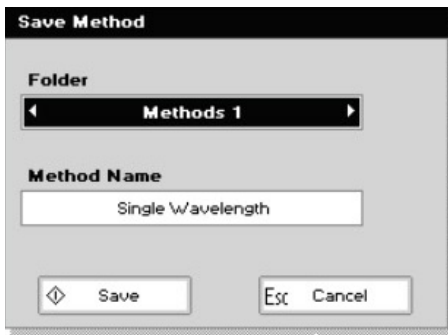
Implement the changes and return to the measurement screen by selecting OK using the confirm key.

OR

Reject the changes and return to the measurement screen by selecting “Cancel” using the escape key

USER INTERFACE

8



Sample Number...

Open save method options.

Select the folder location in which to save the method using the left and right arrow keys.

Press the down arrow, Insert up to a 25-digit method name using the alphanumeric keys.

Save the method and return to the measurement screen by selecting OK using the confirm key.

OR

Return to the measurement screen without saving the method by selecting "Cancel" using the escape key.

9

Auto-Print – Toggle auto-print on or off.

Status Bar Icons

During the measurement process, various status icons are displayed in the top righthand corner status bar. Which icon are displayed depends on the current process being undertaken and the defined settings.



Auto-print to computer via Bluetooth is active.



Auto-print to built-in printer is active.



Auto-print to computer via USB cable is active.



USB flash drive is inserted.



Taking measurement.



Printing to computer via the Bluetooth connection.



Printing to the built-in printer.



Printing to computer via the USB cable connection.



Sending data output to inserted USB flash drive.

USEFUL CALCULATIONS

Beer-Lambert Law

$$A = \epsilon cl$$

A is the absorbance, which although unit-less is usually described as A or AU (absorbance units).

c is the concentration in molar units (M).

ϵ is the molar extinction coefficient in per molar unit per cm (M⁻¹cm⁻¹).

l is the pathlength in centimetres (cm).

As the absorbance value is the known quantity, the Beer-Lambert equation can be rearranged to make concentration (c) the product:

$$c = \frac{A}{\epsilon \times l}$$

Alternative extinction coefficients can be applied to calculate the concentration in alternative units

ϵ	c
Molar extinction coefficient (M ⁻¹ cm ⁻¹)	Molar, or moles per litre, concentration (M, mol L ⁻¹)
Mass extinction coefficient (g ⁻¹ cm ⁻¹)	0.1% Mass per volume concentration (g L ⁻¹)
E1% extinction coefficient (mg ⁻¹ mL ⁻¹ cm ⁻¹)	1% Mass per volume concentration (10 g L ⁻¹)

Conversions between molar, mass, E1% extinction coefficients:

$$\frac{\text{Molar Extinction Coefficient}}{\text{Molecular Weight (g mol}^{-1}\text{)}} = \text{Mass Extinction Coefficient}$$

$$\text{Mass Extinction Coefficient} \times 10 = \text{E1\% Extinction Coefficient}$$

When E1% extinction coefficient are used, the absorbance is multiplied by 10 to present the concentration as a 0.1 % weight per volume (w/v) unit in keeping with convention:

$$c = \frac{A \times 10}{\text{E1\%} \times l}$$

USEFUL CALCULATIONS

Nucleic Acid Concentrations

$$\text{Concentration} = (A_{260} - A_{320}) \times \text{Factor} \times \text{Pathlength Factor} \times \text{Units Factor} \times \text{Dilution Factor}$$

A₂₆₀ is the absorbance at 260 nm.

A₃₂₀ is the optional background absorbance at 320 nm.

Factor is the value defined within the application method parameters.

Pathlength Factor is based on the pathlength selected:

SELECTED PATHLENGTH	PATHLENGTH FACTOR
10 MM	1
5 MM	2
1 MM	10
0.5 MM	20
0.2 MM	50
0.125 MM	80

Units Factor is based on the units selected:

SELECTED UNITS	UNITS FACTOR
MG/ML	1
NG/ML	1
MG/ML	0.001
PMOL/ML ACGT	CALCULATED FROM NUCLEOTIDE SEQUENCE*
PMOL/ML	USER DEFINED+

* Calculated using method described by Ahnert and Patel (197, p. 272), specifically:

$$\text{Molar Extinction Coefficient (M-1cm-1)} = 15\,200 \times \text{Number of A} + 7\,050 \times \text{Number of C} + 12\,010 \times \text{Number of G} + 8\,400 \times \text{Number of T}$$

The units factor is $1\,000\,000 \div \text{Molar Extinction Coefficient (M-1cm-1)}$.

+ The user defined coefficient for pmol/μl has to be $1\,000\,000 \div \text{Molar Extinction Coefficient (M-1cm-1)}$.

Dilution Factor is the value defined within the application method parameters.

USEFUL CALCULATIONS

Protein Concentrations

Concentration = $\frac{((A_{280} - A_{320}) \times F_{280}) - ((A_{260} - A_{320}) \times F_{260})}{\text{Pathlength Factor} \times \text{Units Factor} \times \text{Dilution Factor}}$

A280 is the absorbance at 280 nm.

A320 is the optional background absorbance at 320 nm.

F280 and F260 are the factors associated with the mode selected:

MODE	F280	F260
CHRISTIAN WARBURG	1.55	0.76
BSA	1.49	N/A
IGG	0.73	N/A
LYSOZYME	0.38	N/A
MOLAR EXTINCTION*	$\frac{\text{MOLECULAR WEIGHT}}{\text{MOLAR EXTINCTION}}$	N/A
MASS EXTINCTION*	$\frac{1}{\text{MASS EXTINCTION}}$	N/A
E 1%*	$\frac{10}{\text{E 1\%}}$	N/A
CUSTOM	CUSTOM	CUSTOM

* Molecular weight, molar extinction, mass extinction, and E 1% are the respective values defined within the application method parameters.

Pathlength Factor is based on the pathlength selected:

SELECTED PATHLENGTH	PATHLENGTH FACTOR
10 mm	1
5 mm	2
1 mm	10
0.5 mm	20
0.2 mm	50
0.125 mm	80

Units Factor is based on the units selected:

SELECTED UNITS	UNITS FACTOR
µg/mL	1000
ng/µL	1000
µg/µL	1
mg/mL	1

Dilution Factor is the value defined within the application method parameters.

USEFUL CALCULATIONS

Nucleic Acid and Protein Purity Ratios

$$A260/A280 = \frac{A260 - A320}{A280 - A320}$$

$$A260/A230 = \frac{A260 - A320}{A230 - A320}$$

A260 is the absorbance at 260 nm.

A280 is the absorbance at 280 nm.

A230 is the absorbance at 230 nm.

A320 is the optional background absorbance at 320 nm.

References:

Measuring protein concentration in the presence of nucleic acids by A280/A260: The method of Warburg and Christian. (2006). Cold Spring Harbor Protocols (1).

Fluorescent Dye Quantity

$$\text{Quantity (pmol)} = (A_{\text{dye}} - A320) \times [\text{Pathlength Factor}] \times \text{Volume} \times \text{Dilution Factor} \times \frac{1\,000\,000}{\text{Extinction Coefficient}}$$

A_{dye} is the absorbance value at the dye λ_{max}.

A320 is the optional background absorbance at 320 nm.

Pathlength Factor is based on the pathlength selected:

SELECTED PATHLENGTH	PATHLENGTH FACTOR
10 mm	1
5 mm	2
1 mm	10
0.5 mm	20
0.2 mm	50
0.125 mm	80

Volume is the value defined within the application method parameters.

Dilution Factor is the value defined within the application method parameters.

Extinction Coefficient is the value defined within the application method parameters.

USEFUL CALCULATIONS

Fluorescent Dye Concentration

$$\text{Concentration (pmol/}\mu\text{l)} = \frac{\text{Quantity}}{\text{Volume}}$$

Quantity is the calculated fluorescent dye quantity.

Volume is the value defined within the application method parameters.

Fluorescent Frequency of Incorporation (FOI)

$$\text{FOI (dye/kb)} = \frac{(\text{A}_{\text{dye}} \times 1\,000\,000 \times \text{Molecular Weight})}{(\text{Extinction Coefficient} \times \text{A}_{260} \times \text{Factor})}$$

A_{dye} is the absorbance value at the dye λ_{max} .

Molecular Weight is fixed 324.5 g mol⁻¹ which is an average molecular weight of the nucleotides

Extinction Coefficient is the value defined within the application method parameters.

A_{260} is the dye corrected absorbance at 260 nm.

Factor is the value defined within the application method parameters.

Fluorescent Dye Incorporation

$$\text{Dye Incorporation (pmol/}\mu\text{g)} = \frac{\text{FOI} \times 1\,000}{\text{Molecular Weight}}$$

FOI is the calculated Frequency of Incorporation.

Molecular Weight is fixed 324.5 g mol⁻¹ which is an average molecular weight of the nucleotides

Melting Temperature (T_m)

$$T_m (\text{°C}) = \frac{\Delta H}{(16.6 \times \log_{10}[\text{Buffer}] + (\alpha + \Delta S + (R \times \ln[c \div 4])) - 273.15}$$

ΔH is the change in enthalpy (kcal mol⁻¹) and ΔS is the change in entropy (kcal K⁻¹ mol⁻¹), and are the sum values of their nearest-neighbour pair values, specifically:

USEFUL CALCULATIONS

MOLECULE PAIR*	DNA		RNA	
	ΔH	ΔS	ΔH	ΔS
AA:TT/UU	-9.1	-0.0240	-6.6	-0.0184
AT/AU:TA/UA	-8.6	-0.0239	-5.7	-0.0155
TA/UA:AT/AU	-6.0	-0.0169	-8.1	-0.0226
CA:GT/GU	-5.8	-0.0129	-10.5	-0.0278
GT/GU:CA	-6.5	-0.0173	-10.2	-0.0262
CT/CU:CG	-7.8	-0.0208	-7.6	-0.0192
GA:CT/CU	-5.6	-0.0135	-13.3	-0.0355
CG:GC	-11.9	-0.0278	-8.0	-0.0194
GC:CG	-11.1	-0.0267	-14.2	-0.0349
GG:CC	-11.0	-0.0266	-12.2	-0.0297

* The nucleotide pair on the left of the colon is the 5' to 3' sequence while the nucleotide pair on the right of the colon is the 3' to 5' sequence.

The nucleotide pair on the left of the forward-slash is the DNA sequence while the nucleotide pair on the right of the forward-slash is the RNA sequence.

Buffer is the buffer concentration (M).

α is the helix initiation factor fixed at $-0.0108 \text{ kcal K}^{-1} \text{ mol}^{-1}$.

R is the gas constant fixed at $0.001987 \text{ kcal K}^{-1} \text{ mol}^{-1}$.

c is the calculated nucleic acid concentration (M), specifically:

$$c = A_{260} \times \text{Calculated Factor}$$

A_{260} is the absorbance at 260 nm.

Calculated Factor is calculated from the molecular weight and molar extinction coefficient (E).

$$\text{Calculated Factor} = \frac{\text{Molecular Weight}}{\text{Molar Extinction Coefficient}}$$

Molecular Weight is calculated for the base sequence defined within the application method parameters.

Molecular Weight of DNA = $312.2 \times \text{number of A} + 288.2 \times \text{number of C} + 328.2 \times \text{number of G} + 303.2 \times \text{number of T} + K$

Molecular Weight of RNA = $312.2 \times \text{number of A} + 288.2 \times \text{number of C} + 328.2 \times \text{number of G} + 289.2 \times \text{number of U} + K$

K is phosphorylation constant, specifically

$$K = 17 + (N+2) \times \text{Counter Ion} \\ \text{(Phosphorylated)}$$

$$K = -61 + (N+1) \times \text{Counter Ion} \\ \text{(Non-phosphorylated)}$$

USEFUL CALCULATIONS

N is the base sequence length

Counter ion is the molecular weight of the counter ion selected:

COUNTER ION	PATHLENGTH FACTOR
Na	23.00
K	39.10
TEA1	102.2
TEOA2	149.19
Other*	Custom

¹Triethylamine

²Triethanolamine

*Other is values defined within the application method parameters.

Molar Extinction Coefficient is calculated for the base sequence defined within the application method parameters, and is the sum values of their nearest-neighbour pair values, specifically:

	A	C	G	T/U*
A	13.7	10.6	12.5	11.4
C	10.6	7.3	9.0	7.6
G	12.6	8.8	10.8	10.0
T/U*	11.7/12.3	8.1/8.6	9.5/10.0	8.4/9.8

*The nucleotide on the left of the forward-slash is the DNA sequence while the nucleotide on the right of the forward-slash is the RNA sequence.

OD 600

$$\text{OD} = A_{600} \times \text{Correction}$$
$$\text{Cell/ml} = \frac{A_{600} \times \text{Correction}}{\text{Factor}}$$

A₆₀₀ is the absorbance at 600 nm.

Correction is value defined within the application method parameters.

Factor is the value defined within the application method parameters.

TROUBLESHOOTING

Negative absorbance readings	<ul style="list-style-type: none"> • Sample measurements will be negative absorbance reading if the absorbance value of the reference is higher than the sample. • Negative readings can also result if reference and sample are interchanged or if the sample is very dilute and close to the absorbance of the reference.
Unexpected results	<ul style="list-style-type: none"> • Bubbles or contamination in the sample or reference can result in considerable errors. • Incorrect cuvette orientation. Rotate by 90° and repeat. • Incorrect cuvette material for UV measurement wavelengths. • Wrong pathlength selected in software. • Contact your supplier for advice on the minimum concentrations that can be measured.
Absorbance higher than expected	<ul style="list-style-type: none"> • Incorrect sample reference. • Incorrect cuvette orientation. • Incorrect cuvette material for measurement wavelengths. • Wrong pathlength selected in software. • Contamination in sample or on cuvette. • Check baseline, if greater than 0 A toggle background correction or us an appropriate reference sample. • Possible incorrect optical alignment. Contact technical support.
Absorbance lower than expected	<ul style="list-style-type: none"> • Incorrect sample reference. • Check sample and reference for contamination. • Check sample and reference samples are not the same. • Incorrect cuvette material for measurement wavelengths. • Wrong pathlength selected in software. • Check the beam height and buffer sample volume. • Check baseline, if greater than 0 A toggle background correction or us an appropriate reference sample. • Possible stray light issue. Contact technical support.
Poor reproducibility	<ul style="list-style-type: none"> • Insufficient sample in cuvette. • Cuvette in wrong orientation. • Cuvette material unsuitable for wavelengths used. • Concentration of sample too low or too high. For best results, the measured sample absorbance using a 10 mm pathlength cuvette should ideally be between 0.1 and 1.0 A. If absorbance is >1 A, measurement is no longer in the most linear range. • Particulates in sample. Absorbance measurements will not be accurate with turbid samples. • Possible noise or measurement stability issue. Contact technical support
Instrument start up reported failure	<ul style="list-style-type: none"> • Check the cell holder is empty. • Check original 18V dc supply is connected and is fully engaged. • Report persistent failures to technical support.

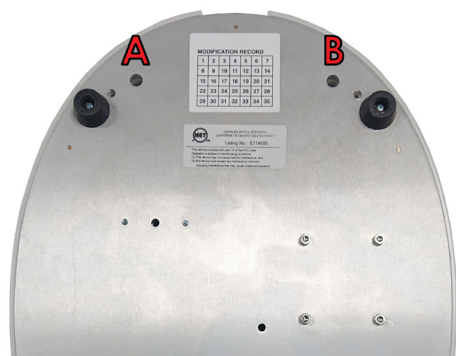
BUILT-IN PRINTER

Built-in Printer Accessory Part Numbers

80-3003-84 – Built-in Printer accessory

80-3004-07 – Spare paper for printer (20 rolls)

Printer Installation Guide



Step 1

Invert the instrument and remove the screws from positions A and B.



Step 2

Return the instrument to its upright position and lift the accessory cover vertically upwards.



Step 3

Remove any tie-wraps from the printer cable and insert its connector to the socket on the underside of the printer.

BUILT-IN PRINTER



Step 4

Lower the printer onto the locating bosses and push down firmly.



Step 5

Invert the instrument and replace the accessory cover screws at A and B.

Refilling the Printer Paper



Step 1

Lift off the printer cover using the tabs and discard the spent roll.



Step 2

Open the platen lock, feed in the paper so that the loose end is underneath the roll pointing towards the front of the instrument, then close the platen lock and draw the loose end through using the platen wheel.

BUILT-IN PRINTER



Step 3

Replace the printer cover so that the loose end sits above it.

Ordering Information

Part Number	Description
80-3007-32	WPA Biowave 3
80-3007-33	WPA Biowave 3 with Printer
80-3007-34	WPA Biowave 3 with Bluetooth
80-3007-35	WPA Biowave 3 with Printer and Bluetooth
80-3007-37	WPA Biowave 3+
80-3007-38	WPA Biowave 3+ with Printer
80-3007-39	WPA Biowave 3+ with Bluetooth
80-3007-40	WPA Biowave 3+ with Printer and Bluetooth

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