OPTIONAL SOFTWARE - E-SEF Enabled

SM1 Nucleic Acid and Protein - more advanced calculations:

Warburg and Christian	
DNA	260/280
DNA	260/280, 320
Protein	260/230
Protein	260/230, 320

260/280
260/280, 320
260/230
260/230, 320

- SM2 Molecular calculations Molecular concentrations Recovery of oligonucleotides Phosphate concentration Protein purity Theoretical thermal melt
- SM3 Protein Assay Bradford 595nm
- SM4 Bacterial Growth Optical density at 600nm
- SM5 Validation: Wavelength, Absorbance, Bandwidth etc
- SM6 Method Storage for 30 methods security protected

ACCESSORIES AND SPARES

Cell holder for 50µL cells - type E	2020 39 00
Printer connection cable	8000 71 00
Dot matrix printer	8000 70 01
Spare deuterium lamp	2202 01 42
2 Certified wavelength filters with certificate	303 40 00
4 Certified Absorbance filters with certificate	594 99 00
Dust cover	2020 01 23

SPECIFICATION

Wavelength Range	190-370nm, 595 and 600nm
Wavelength Accuracy	±1nm
Wavelength Reproducibility	±0.1nm
Wavelength Calibration	Automatic at switch on
Wavelength Selection	Automatic for assay selected
Monochromator	Littrow 1200 lines per mm holographic grating
Optical Bandwidth	4nm
Straylight	Less than 0.02% at 280nm (Acetone)
Photometric Range	-0.3 – 3 Absorbance
Photometric Linearity	±1% or ±0.005, whichever is greater
Stability	Better than 0.001A/hour at 240nm constant T.
Real Time Clock	Timed and dated reports
Light Source	Deuterium lamp with life indicator
Printer Output	Centronics parallel port
Power Requirements	110-240V, 50/60Hz, 100W
Size and Weight	480 x 340 x 205, 17.6Kg, (18.5kg printer version)

GeneQuest™ is a registered Trade Mark of Cecil Instruments Ltd. Cecil Instruments policy is one of continuous development. We therefore reserve the right to change specification without notice.

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BENEFITS

Full screen display of menus, prompts and results

Software options password enabled at any time by E-SEF

Storage for 30 user methods, password protected

Warning of poor sample transmission

Signal enhancement for poor sample transmission

Plot of curve fitted to standards

Full UV range available to create methods

ORDERING

Each GeneQuest analyser is supplied complete with an ultra microcell holder, power cable, operator's manual and short form operating instructions.

CE 2301 Analyser 190-370nm, 595 and 600nm 4nm optical bandwidth

CE 2302 Analyser 190-370nm, 595 and 600nm 4nm optical bandwidth With integral printer

ISO 9000

CERTIFIED COMPANY













Generouest™ dedicated to dna / RNA



Dedicated to Molecular Biology

The GeneQuest[™] Analyser is a spectrophotometer dedicated to the requirements of the Molecular Biology Laboratory. Fully programmed methods include those for RNA, ds DNA, ss DNA, oligonucleotide primer, proteins, bacterial growth and cell culture measurements and molecular calculations, including theoretical thermal melt.

Ultra Micro-Samples

Measurements are possible for a wide range of sample volumes, from 2000μ L to 5μ L using an appropriately selected cell in the ultra microcell holder fitted as standard in the GeneQuestTM.

STANDARD SOFTWARE

- A260/A280 Ratio
- A260/A280 with background correction at 320nm
- A260/A230 Ratio
- A260/A230 with background correction at 320nm

Simplicity of Operation

Insert the reference, press ZERO, then insert the sample and press RUN to obtain a fully documented report complete with header, sample number and space for the operators name etc. Operation could not be more simple.

Operators may create and store up to 30 of their own methods with the method storage option.

Automatic Wavelength Selection

All the fully programmed methods are selected from a menu and displayed with headers, prompts and details on the large display screen on which information may be scrolled for viewing.

All wavelengths for any given assay are automatically selected and measured. Results are displayed on the screen and a timed and dated printout is available on an external printer, or the internal printer of the CE2302.

Enhanced Performance

The excellent accuracy of the GeneQuest[™] for normal samples is further enhanced for poorly transmitting samples by an automatic integration process initiated in response to an alert message.

Validation

To fulfil legislative requirements and GLP, onboard software allows for performance validation using filter sets calibrated by Cecil Instruments and supplied with calibration certificates traceable to NPL standards. Methods for liquid samples are also provided.

- ss DNA concentration
- ds DNA concentration
- RNA concentration

D

Direct UV method for protein at 280nm

A range of UV transmitting silica cells is available. The listing given here enables the cell appropriate to the sample volume to be chosen. All cells except the standard must be of the self masking type.

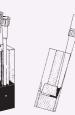
Using a 10mm pathlength cell an absorbance value of 1 is estimated for a concentration of $50\mu L/mL$ of ds DNA



Volume (µL)	Pathlength (mm)	Cell Description	Cell Type	Part Number
2000	10	Standard	A	202 07 26
500	10	Semi-micro, thick base	В	202 07 57
250	10	Micro, thick base	С	202 07 38
70	10	Ultra micro	D	202 07 15
50	10	Ultra micro	Е	2020 07 01
		use holder 2020 39 00		
10	10	Ultra micro, pipette filling	F	202 07 64
1000	5	Standard - use spacer	A	202 07 25
5	5	Ultra micro, pipette filling	F	202 07 63







AND MOLECULAR BIOLOGY

D2	260	260.0		0.836.		
Mea	surement	finish	led	Rat	io 260/280	
One	rator:	JN				
	erence:					
	ple:		3			
	elength					
	= readin					
	= readin					
	io = A1					
IRa	tio = A2	/A1				
• • •						
ADS	orbance		n-+	TRatia		
		length 280.0	Racio	IRACIO		
		280.0				
	ple 0.459	0 457	1 004	0 006		
	0.459					
4	0.807	0./94	1.010	0.284		

0.321 0.313 1.026 0.975

^{D2} 280.0 _{nm}	0.000
	Theoretical Tm
THEORETICAL TM CECIL CE 2302 Serial No: 141099 S/W version: T0231 Time: 09:08 28/10/99 Molarity = 0.050 mol/L MerLength = 54 %GC = 42 % Tm = 81.5 + 16.6*log(Mol 675/MerLength = 65 'C	larity) + 0.41*%GC -

	_	
	· · · · · · · · · · · · · · · · · · ·	
		1

^{D2} 280.0 _{nm}	7.325 _{mg/l}
	Direct UV
DIRECT UV CECIL CE 2302	
Serial No: 141099 S/W version: T0231	
Time: 09:01 28/10/99 Bandwidth: 4.0 nm Conc curve: quadratic	
Stand Dev: 0.00 Corr coeff: 1.0000 Wavelength: 280.0 nm	
Std Abs Std Calc No. conc conc	
2 1.188 26.00 26.00 1 0.794 18.00 18.00	

QUANT MENU
CECIL CE 2302
Serial No: 141099
S/W version: T0231
Time: 10:09 28/10/99
Bio Analysis
SSDNA
dsDNA RNA
DNA Ratio Ratio 260/280
Ratio 260/280 Ratio 260/280,320
Ratio 260/280,320 Ratio 260/230
Ratio 260/230 Ratio 260/230,320
Warburg & Christian
DNA 260/280
DNA 260/280,320
Protein 260/280
Protein 260/280,320
Kalb & Bernlohr
DNA 260/230
DNA 260/230,320
Protein 260/230
Protein 260/230,320
Theoretical Tm
Tm (Wallace)
Tm (Baldino)
Bacterial Growth
Protein Assay
Bradford
Direct UV

		DFO		
	CECIL			,
`				
Ser:	ial No	:	1410	99
s/w	versi	on:	T023	1
Time	∋: 08	:53	28/1	0/99
Band	dwidth	:	4.0	nm
Cond	curv	e:	line	ear
Star	nd Dev	:	0.06	;
Cori	c coef	f:	0.99	99
Wave	elengt	h:	595.	0 nm
· • ·	cator:			
Stai	idard:			
Std	Abs	SI	.d	Calc
No.		c	onc	conc
3	0.931	18	3.00	18.03
2	0.556	1:	L.00	10.92
1	0.353	7	000	7.053
a	: = +0	250	10	0.0+3
lond	2 = +0	. 350) +19	.00*A

Assays Displayed on Screen

Shown here are three screen displays of typical assays, for protein by the direct UV method, ratios at 260/280nm and a theoretical melt Tm calculation.

Printout of Assays

The printouts shown here are for typical assays using the Bradford method at 595nm, the determination of ds DNA, and the results of the Kalb Bernlohr determination of nucleic acids and proteins.

A plot of any curve or straight line fitted to a series of standards is always available to verify the fit.

The printout on the left of this page shows a list of all the available methods offered by the GeneQuest[™].

DSDNA				
CECIL CE	2302			
Serial No:	141099			
S/W version:	T0231			
Time: 15:43	27/10/99			
Bandwidth:	4.0 nm			
Wavelength:	260.0 nm			
Operator:	JN			
Reference:	Air			
Sample:	RS/553			
Wavelength for	ormula:			
A1 = reading	at 260.0	nm		
Dilution =	2			
PathLength =	10 mm			
Factor =				
Conc =	A1 * Fact	or * 10 * Di	lution / PathLeng	th
ug/mLdsDNA				
Avogadro =		3		
MolWt =				
			olWt*1e6) /mL	
MolConc =	Conc * 1e	.3 / MolWt pm	nol/uL	
Absorbance	Cons	Molecules	MolCong	
	g/mLdsDNA		pmol/uL	
Sample	3/ IIIIGSDNA	/ 1111	pmor/un	
Sampie 1 0.319	31 60	1.213e+17	201.4	
		1.741e+17		
		3.068e+17		

0.461 46.10 1.753e+17 291.0

5

CECIL CE 2302 Serial No: 141099 S/W version: T0231 Time: 10:27 28/10/99 Bandwidth: 4.0 nm Path length: 10 mm Operator: JN Reference: Air Sample: WQ/740 Wavelength formula: Al = reading at 230.0 nm
S/W version: T0231 Time: 10:27 28/10/99 Bandwidth: 4.0 nm Path length: 10 mm Operator: JN Reference: Air Sample: WQ/740 Wavelength formula:
Time: 10:27 28/10/99 Bandwidth: 4.0 nm Path length: 10 mm Operator: JN Reference: Air Sample: WQ/740 Wavelength formula:
Bandwidth: 4.0 nm Path length: 10 mm Operator: JN Reference: Air Sample: WQ/740 Wavelength formula:
Path length: 10 mm Operator: JN Reference: Air Sample: WQ/740 Wavelength formula:
Operator: JN Reference: Air Sample: WQ/740 Wavelength formula:
Sample: WQ/740 Wavelength formula:
Wavelength formula:
A1 = reading at 230.0 nm
A2 = reading at 260.0 nm
A3 = reading at 320.0 nm
Dilution = 4 Conc = ((A2-A3)*49.1 - (A1-A3)*3.48) * Dilution ug/mLDNA
Avogadro = 6.02214e23
MolWt = 213.7 q
Molecules = Conc * Avoqadro / (MolWt*1e6) /mL
MolConc = Conc * 1e3 / MolWt pmol/uL
ExMolConc = 1150 pmol/uL
Recovery = 100 * MolConc / ExMolConc %
PhosConc = Conc/315.0 pmol/uLP04
Absorbance A
Wavelength Conc Molecules MolConc Recovery PhosConc 230.0 260.0 320.0 ug/mLDNA /mL pmol/uL % pmol/uLPO4
Sample
1 0.335 0.320 0.305 2.528 7.125e+15 11.83 1.029 0.008
2 0.336 0.320 0.306 2.332 6.572e+15 10.91 0.949 0.007
3 0.338 0.322 0.307 2.514 7.086e+15 11.77 1.023 0.008