UV/VIS SPECTROPHOTOMETER USER'S MANUAL

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Contents

Safety:

The safety statements in this manual comply with the requirements of the HEALTH AND SAFETY AT WORK ACT, 1974.

Read the following before installing and using the instrument and its accessories. This instrument should be operated by appropriate laboratory technicians.

General:

The apparatus described in this manual is designed to be used by properly trained personnel in a suitable equipped laboratory. For the correct and safe use of this apparatus it is essential that laboratory personnel follow generally accepted safe procedures in addition to the safety precautions called for in this manual.

The covers on this instrument may be removed for servicing. However, the inside of the power supply unit is a hazardous area and its cover should not be removed under any circumstances. There are no serviceable components inside this power supply unit. For this instrument, avoid touching the high voltage power supply at all times.

Some of the chemicals used in spectrophotometry are corrosive and/or inflammable and samples may be radioactive, toxic, or potentially infective. Care should be taken to follow the normal laboratory procedures for handling chemicals and samples.

Electrical:

Before switching on the apparatus, make sure it is set to the voltage of the local power supply (see Installation).

The power cord shall be inserted in a socket provided with a protective earth contact. The protective action must not be negated by the use of an extension cord without a protective conductor.

Warning:

Any interruption of the protective conductor inside or outside the apparatus or disconnection of the protective earth terminal is likely to make the apparatus dangerous. Intentional interruption is prohibited.

Whenever it is likely that the protection has been impaired, the apparatus shall be made inoperative and be secured against any unintended operation.

Note: NEVER touch or handle the power supply on this instrument due to the high voltage!

The protection is likely to be impaired if, for example, the apparatus

- Shows visible damage
- Fails to perform the intended measurements
- · Has been subjected to prolonged storage under unfavorable conditions
- Has been subjected to severe transport stresses

Performance:

To ensure that the instrument is working within its specification, especially when making measurements of an important nature, carry out performance checks with particular reference to wavelength and absorbance accuracy. Performance checks are detailed in this manual.

Radio Interference:

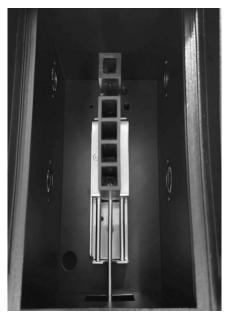
For compliance with the EMC standards referred to in the EC Declaration of Conformity, it is necessary that only shielded cables supplied by us are used when connecting the instrument to computers and accessories.

Introduction:

This instrument (Fig 1) is a double beam, general purpose instrument designed to meet the needs of the Conventional Laboratory, This instrument is ideal for various applications, such as: Chemistry, Biochemistry, petrochemistry, Environmental Protection, Food and Beverage Labs, Water and Waste Water Labs and other fields of quality control and research.

This instrument incorporates a 320×240 dot matrix LCD display for photometric results, easy operation and wavelength range of 190nm to 1100nm. This instrument is ideal for measurements in the visible and ultraviolet wavelength region of the electromagnetic spectrum.





The Sample Compartment



Fig1

Working Principle:

The spectrophotometer consists of five parts: 1) Halogen or deuterium lamps to supply the light; 2) A Monochromator to isolate the wavelength of interest and eliminate the unwanted second order radiation; 3) A sample compartment to accommodate the sample solution; 4) Two detectors to receive the transmitted light and convert it to an electrical signal; and 5) A digital display to indicate absorbance or transmittance. The block diagram (Fig 2) below illustrates the relationship between these parts.

Block diagram for the Spectrophotometer

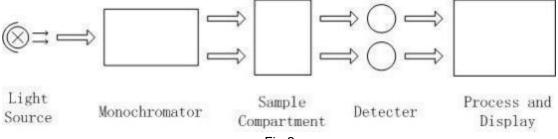


Fig 2

In your spectrophotometer, light from the lamp is focused on the entrance slit of the monochromator where the collimating mirror directs the beam onto the grating. The grating disperses the light beam to produce the spectrum, a portion of which is focused on the exit slit of the monochromator by a collimating mirror. From here the beam is passed to a sample compartment through one of the filters, which helps to eliminate unwanted second order radiation from the diffraction grating. Upon leaving the sample compartment, the beam is passed to the silicon photodiode detector and causes the detector to produce an electrical signal that is displayed on the digital display.

Unpacking Instructions:

Carefully unpack the contents and check the materials against the following packing list to ensure that you have received everything in good condition.

	Packing List	
	Description	Quantity
•	Spectrophotometer	1
•	Mains Lead	1
•	Glass Cuvettes	1 Set of 4
•	Quartz Cuvettes	1 Sets of 2
•	Dust Cover	1
•	Operation Manual	1
•	Software Manual	1
•	Software Kit (Disc 1+ USB Lead 1+ Dongle 1)	1

Note: The printer and auto-cell holder mentioned in this manual are all optional accessories, they do not come standard with the instrument.

Specifications:

Spectral Bandpass	2nm	
The Optical System	1200/mm Diffraction Grating	
Wavelength Range	190-1100nm	
Wavelength Accuracy	±0.5 nm	
Wavelength Repeatability	< 0.2nm	
Stray Light	≤0.05%@220nm&360nm	
Photometric Range	0-200%T, -4-4.0A, 0-9999C (0-9999F)	
Stability	± 0.0015A @ 500nm	
Power Requirements	AC 110V/60Hz or 220V/50Hz	
Data Output Port	USB mouth, Centronics	
Dimension (L*W*H)	610*530*360mm	
Net Weight	25kg	

Installation:

- 1. After carefully unpacking the contents, check the materials with the packing list (page 4) to ensure that you have received everything in good condition.
- 2. Place the instrument in a suitable location away from direct sunlight. In order to have the best performance from your instrument, keep it as far as possible from any strong magnetic or electrical fields or any electrical device that may generate high-frequency fields. Set the unit up in an area that is free of dust, corrosive gases and strong vibrations.
- 3. Remove any obstructions or materials that could hinder the flow of air under and around the instrument.
- 4. Use the appropriate power cord and plug into a grounded outlet.
- 5. Turn on your spectrophotometer. Allow it to warm up for 15 minutes before taking any readings. We suggest you then do the Calibrate System with the Search 656.1nm to set the wavelength to the deuterium lamp emission line.

NOTE:



This symbol means Caution, Risk of Danger.

Operation:

Prepare the spectrophotometer

Fig 3 is the control panel. User can perform all operations by pressing the keys and all the results and operation information are displayed on the LCD.



Fig 3

Description of keys

[LOAD]	Load data or curve saved before;
[SAVE]	Save data or curve;
[SET]	Set wavelength;
【ZERO】	Blank or scan the user base line;
[PRINT]	Print test results or screen
[START]	Start testing or scanning sample;
[ESC/STOP]	Exit to previous screen or cancel the operation;
[ENTER]	Confirm the inputted data or selected item; Go into next
	setup or screen;
【F1】-【F4】	Function based on the information on the screen;
【0】-【9】	Input number or letter, consecutively press a numeric key
	to select a character;
【+/-/.】	Input +,- or dot;
【CE】	Clear all characters when you are inputting or clear curve
	displays on the screen;

【<】,【>】	Change "x" scale; Search point after scan; 【<】 clear a character;
【∧】,【∨】	Change "y" scale; Search peak after scan; Scroll items for selecting; Change capital/small letter last typed in; Browse the items for selection;
[CELL]	Set cell position (Only available when Auto Changer used).
(HELP)	Reserved key for future Function Extending, not available now.

<u>Turn on spectrophotometer</u>

Turn on spectrophotometer by pressing the Power Switch (IO)(see Fig1). The instrument starts to initiate and the steps are as below:

1. The instrument will check memory first (Fig 4), please wait or press any key to skip this step ,after positioning filter, auto-cell changer(if installed) and D2/Wlamps, the screen display as Fig 4A. 15 minutes pass or press [ESC], the screen display as Fig 5,Select "No" to skip to main menu(Fig 7) and select "Yes"(recommended) to calibrate system (Fig 6). The calibrating process include "get dark current", "searching 656. 1nm" and "check energy". After finish the calibration system, go to main menu too (Fig 7).

2. If the data in memory has been lost, the instrument will directly calibrate system without any choice for you.

state system without any choice for you.

WL: 656.2nm	RAM Checked: 16kb
Wait until EasyRTOS booted: Check memory	D2∋ W ∋
UV/Vis Spectrophotometer	
Fig 4	

3. If no auto-cell changer installed "cell #1" will disappear in Fig7



Fig 4A

WL: 656.Inm	 16:20:18
Wait until EasyRTOS booted: Check memory	D2页 W 司
System calibration? 🚾	



WL : 656.1nm	1648:03
Wait until EasyRTOS booted: Check memory	D2 ⊧1 W ∋termen
UV/Vis Spectrophotometer	

Fig 6

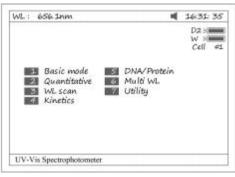


Fig 7

Basic operation

※ Blank

There is a system baseline stored in the memory of the instrument. Usually user may not rebuild system baseline before test. Only putting the sample into the sample light path and the reference into the Reference Light Path, the result can be obtained. As the system baseline always get a little change after the instrument is powered on, it is necessary for the user to rebuild the system baseline. There are a couple of ways to rebuild the system baseline. Select "Yes" in Fig 5 or Press **[0]** In Fig 73 or Press **[F4]** In Fig41,

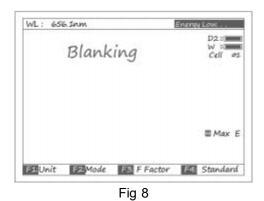
Regarding Blanking, important points list below:

A. Take measure in Basic Mode

 a. Put the reference cuvette with *reference solution* into the Reference Light Path and the sample cuvette with *reference solution* into the Sample Light Path. Press the key 【ZERO】 for blanking.

Note 1. If the reference solution is too thick, "Energy Low..." will appear following the "Blanking..." on the screen (Fig 8). If "Energy too Low..." appears following the "Blanking...", the test will be paused and "Warning..." will appear on the screen. (Fig 9).

2. If no automatic changer installed "cell #1" and "Max E" will disappear in Fig8



3. DO NOT OPEN SAMPLE COMPARTMENT LID DURING BLANKING.

4. The dark current don't be taken after power on if you bypass the calibrating system. It is recommended to take the dark current after warm up.(See page 38).



Fig 9

b. Take out the sample cuvette, replace the *reference solution* with *sample solution* after flushing the cuvette completely.Put the sample cuvette into the Sample Light Path, then the result will display on the screen automatically. However the **[START]** must be pressed in other measurements such as DNA/Protein, Muli WL and Quantitative etc.

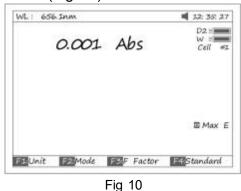
B. Take measure in WL Scan

a. After all scan parameters are entered, put the reference cuvette with *reference solution* into the Reference Light Path and the sample cuvette with *sample solution* into the sample light path,Press **[START]** to scan.

b. (Recommended) After all scan parameters are entered, put the reference cuvette with *reference solution* into the Reference Light Path and the sample cuvette with *reference solution* into the Sample Light Path, Press

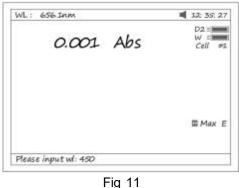
[ZERO] to obtain the user baseline. Then take out the sample cuvette, replace the *reference solution* with *sample solution* after flushing the cuvette completely. Put the sample cuvette into the Sample Light Path. Press **[START]** to scan.

Set wavelength (Example: set wavelength in "Basic mode")

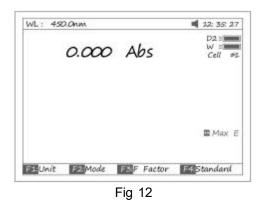


• Press **[SET** λ] (Fig 10).

♦ Use numeric keypad to input wavelength (Fig 11).



Press **[ENTER]** to change the wavelength from 656.1nm to 450.0nm,and then blank; After blanking, the screen displays as Fig 12.



- Load or delete data or curve (Take the "WL scan" test For example) Press 【3】 in Fig.7 go into "WL scan". After 【LOAD】 being pressed, the first file (ABC.wav)in memory will appear on the bottom line of screen .Showed as Fig 13. Press 【八】 or 【\\] to browse the files stroed in memory. Then if :
 - 1. The key **[ENTER]** be pressed, the file selected will be loaded and displays on the screen. Fig 14.
 - Note (1) The file selected must match "WL scan" test's type. If not, the "file type error..." will appear on the Right of top line.
 (2) Different test has different file type. Refer to table 1 on Page 12.
 - 2. The key **[CE]** be pressed, the file selected will be deleted by selecting "Yes".

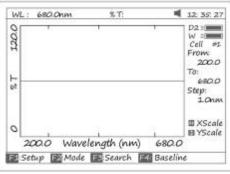


Fig 13

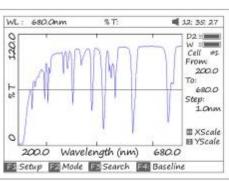


Fig 14

Table 1			
Test	File Type		
Quantitative Curve	*** fit		
Quantitative Test Result	***.qua		
WL Scan	*** wav		
Kinetics	***.kin		
DNA/Protein	***.dna		
Multi WL	*** mul		
WL Validity	***. wlv		
Accu. Validity	***.phv		

% Save data or curve (Example: Save curve in "WL scan")

- Press the key [SAVE] in Fig14 to save curve.
- Name the curve by pressing the numeric keypad (Fig 15), press the key [ENTER] to confirm.
 - Note(1). Pressing numeric key continually to scroll characters and pressing 【 ∧ 】, 【 ∨ 】 to alter capital letter to miniscule. Table 2 shows all characters built in.
 - (2) If the name already exists in memory, the warning "duplicated name, are you sure ? " will appear . "Yes" for overwrite and "No" for Exit.
 - (3) The length of filename is less than 4.

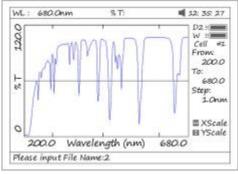


Fig 15

			Table 2		
key	representing	key	representing	key	representing
0	0,+,-,* ,/	1	1,#,?,:,I	2	2,A,B,C,=
3	3,D,E,F,%	4	4,G,H,I,{	5	5,J,K,L,}
6	6,M,N,O,~	7	7,P,Q,R,S,	8	8,T,U,V,"
9	9,W,X,Y,Z	+/-/.	-		

Table 2

※ Print test report (For example: Print the report in "Basic mode", Fig16) Press the key **[PRINT]** to print the report (curve or data you have loaded or tested, Fig 17).



Fig 16

Basic Mode Test Report

Wavelength: 546.0nm Result: 0.221 Abs Date and Time: 25-06-2003 13:55:53

Fig 17

Before measurement

• Make a blank reference solution by filling a clean cuvette (or test tube) half full with distilled or de-ionized water or other specified solvent. Wipe the cuvette with tissue to remove the fingerprints and droplets of liquid.

Analyze Sample

For different user requirements, we have provided different test methods.

Basic Mode

Push the blank cuvette into the Reference Light Path and Main Light Path. In main menu (Fig7),press **[1]** to enter "Basic mode" test . After automatically blanking, it will display as Fig 18 (automatic changer installed) or Fig 19 (automatic changer uninstalled) and wait for the operator. **[ESC/STOP]** to exit.

Note: If no automatic changer installed "cell #1" and "Max E" will disappear in Fig18



Fig18



Fig19

Contract Contract

There are three modes (T%, Abs, conc/factor) for you to select by pressing **[F2]** to make choice.



Fig 20

1. Abs mode

Push the blank cuvette into the Reference Light Path and Main Light Path. Press **[F2]** to select Abs mode ,Press **[ZERO]** for Blanking , and then Push the sample into Main Light Path to take reading(Fig 20)

2. T% mode

The operation is the same as Abs test mode but pressing \pressing to select T% mode .

3. Conc/Factor mode

Press **[F1]** to select a concentration unit (Fig 21). If no unit is suitable for your test, please select the item "Other", press **[ENTER]** and input a new unit by pressing the numeric keypad (Fig 22).

0.000	mg/ml	D2 ≥ W = Cell ≠2
F factor	2.000	
		⊞ Max E
Please select unit 👔		

Fig 21

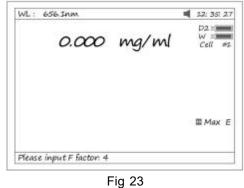
WL: 656.Inm	1 22: 35: 27
0.000 mg/ml	D2 ≡ W ≡ Cell ≠2
F factor 2.000	
	⊞ Max E
Please input self defined unit: 🛙	

Fig 22

- Push the blank cuvette into the Reference Light Path and Main Light Path and press [ZERO] for Blanking. There are now two choices for you to take:
- 4.1 Press **[F3]** to input known F value, Fig 23. Then push the sample into Main Light Path to take reading of concentration
- 4.2 Push sample of known concentration into the Main Light PathPress [F4] to input known Conc value, Fig 24. Then push the sample into Main Light Path to take reading of concentration.

Note:1. You can select wavelength at any time by pressing **[SET** λ **]**. After your selection, instrument always blanks automatically.

2.IfF value is more than 9999,the clue of "out of range" will display on screen.



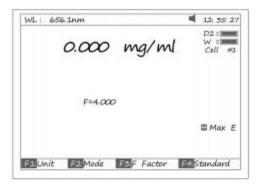


Fig 24

Print Test Report

Press **[PRINT]** to print test results (Fig 25).

Basic Mode Test Report

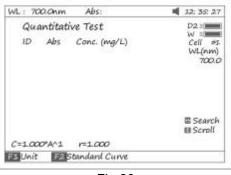
Wavelength: 546.0nm Result: 0.221 Abs Date and Time: 25-06-2003 13:55:53

Fig 25

Quantitative

Press **[2]** in Main Menu for "Quantitative" Test (Fig 26). Press **[ESC/STOP]** to exit.

Note: .If no automatic changer installed "cell #1" will disappear in Fig26.





How to operate

1. Press **[F1]** to select unit of concentration (Fig 27).

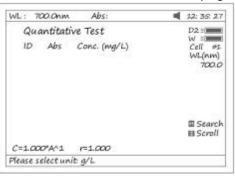


Fig 27

2. Press [SET λ] to select correction methods and enter the wavelength. There are three correction methods (single, Isoabsorbance and 3 point, Fig 28)

Note: Please refer to the Appendix B for the correction method.

WL: 700.0nm	Abs:	12: 35: 27
Quantitat	ive Test	D2 =
ID Abs	Conc. (mg/L)	W =1 Cell ≠5 WL(nm) 700.0
		II Search III Seroll
C=1.000*A^1	r=1.000	
Correction meth	od Single WL	

Fig 28

3. Press **[F2]** in Fig 26 for more items to select .See Fig 29.



Fig 29

- **3.1** Press **[F1]** in Fig 29 to select fitting method. There are 4 methods for you to choose: Linear fit, linear fit through zero, square fit and cubic fit.
- **3.2**Press **[F2]** in Fig 29 to enter directly a known standard curve.Fig29A.

WL: 700.0nm Abs:	12: 35: 27
Calibration table	D2 ±0
No Conc. (mg/L) Abs	Cell == WL(nm) 700.0
C=1.000*A^1 r=1.000 InputK1=1.033	

Fig 29A

The constants to be entered depend on which fitting method selected. The table below lists their relation:

Fitting Method	Fitting Equation	constants
linear fit through zero	C=K1×A	K1, r*
Linear fit	C=K0+K1×A	K0,K1,r*
square fit	$C=K0+K1 \times A+K2 \times A^2$	K0,K1,K2
cubic fit	$C=K0+K1\times A+K2\times A^2+K3\times A^3$	K0,K1,K2,K3

* r : regression co-efficients, default=1

3.3Press **[F3]** in Fig 29 to establish a standard curve by measuring a group of standard samples. See Fig 30.

3.3.1 Enter standard concentrations of samples by pressing the Numeric keypad followed by 【ENTER】. Press 【 ∧ 】 or 【 ∨ 】 to modify the inputted data (Fig31).

Press **[ESC/STOP]** to finish inputting and to exit (Fig 32).

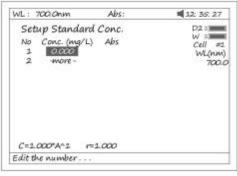


Fig 30

WL:	700.0nm	Abs:	💐 12: 3 <i>5</i> : 27
Se	tup Standan	d Conc.	D2 =
No 1 2 3	Conc. (mg/L 2.000 3:000 -more-) Abs	ν =1 Cell ≠1. WL(nm) 700.0
	.000°A^1 r Standard Con	1000 mm	

Fig 31

3.3.2 Push the blank cuvette into the Reference Light Path and Main Light Path, press **[0Abs/%100T]**, the instrument will step to the wavelength and blank. See Fig 32.

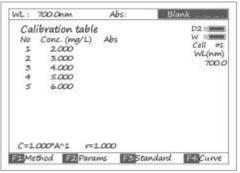


Fig 32

- **3.3.3** Pull the first sample cuvette of known concentration into the light path, Press the key **[START]** to get values of standard curve one by one (Fig 33).
- 3.3.4 Press [F4] to draw the curve. You can get a different curve by pressing [F1] to select a different fitting method. (See Fig 34-Fig37.) For linear fits, "r" represent fitting coefficient of linear regression .r=1 is best fitting.usually "r" is very close to 1.

Note: If there are few standard samples, it is not suitable for selecting square fitting, especially cubic fitting, otherwise invalid fitting result will be obtained.

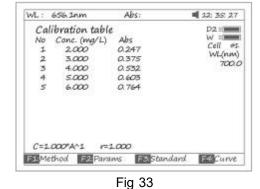


Fig 34 linear through zero fit

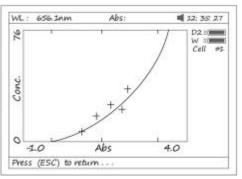


Fig 35 square fit

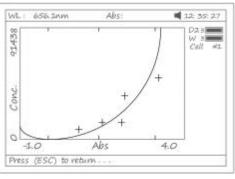


Fig 36 cubic fit

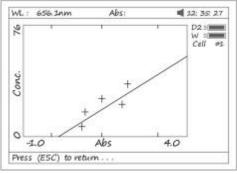
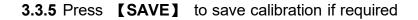


Fig 37 linear fit



3.3.6 Press [ESC/STOP] to exit

4. Quantitative Test

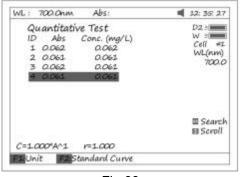
Before test, the standard curve must be obtained. There are three ways for you to obtained it (a, b or c).

- a) Standard curve built up and saved in the instrument.
 In Fig 33 press 【Load 】 and then press 【 ∧ 】 or 【 ∨ 】 to select the file with type ***.fit. At last press 【ENTER】 TO comfirm.
- b) Known standard curve, which is not saved in the instrument. See 3.2. For Fig 29 enter a known standard curve directly.
- c) Use the standard samples for the test. First the standard curve must be established using the method shown in 3.3.

Note: All sample results must be taken in screen Fig26.

4.1 Push the blank cuvette into the Reference Light Path and Main Light Path and press **[ZERO]** for blanking.

4.2 Pull the sample cuvette into Main Light Path, press the key **[START]**, the results will be displayed on the screen (Fig 38).





4.3 If there is more than one sample, repeat step 4.2 for the next sample

4.4 Press (SAVE) to save the results and fitting parameters

Print Test Report

Press the key **[PRINT]** to print the test report (Fig 39). Quantitative Test Report

File Name: Date and Time: 25-06-2003 13:54:32 546.0nm Abs(eff) C(mg/L) No 0.212 0.212 3.315 1 2 0.212 0.212 3.321 3 0.000 0.212 3.315 Fitting Params:C= 15.64*A^1 r= 0.105 Fig 40

WL Scan

Press [3] in main menu for "WL Scan" test (Fig 41). [ESC/STOP] to exit.

To load a previous curve, press **[LOAD]** and select a previously stored curve (.wav)

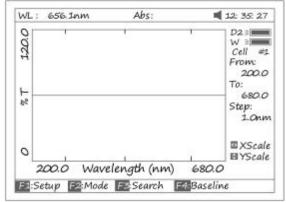
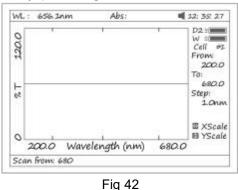


Fig 41

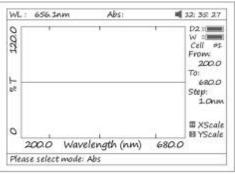
Scan sample

1. Press **[F1]** to setup, input the start wavelength, and end wavelength by pressing the numeric keypad (Fig 42). Note: This instrument scans from high to low wavelength. Browse and select the items of scan step and scan speed by pressing **[** Λ **]** or **[** \vee **]**.



Note: "Scan step" allows the selection of 0. 1nm, 0.2nm ,0.5nm, 1nm ,2nm and 5nm. "Scan speed" allows the selection of "HI", "MEDIUM" and "LOW". For survey scan we suggest 5nm, HI. For detailed scan we suggest 0.5nm, HI

2. Press **[F2]** to select the test mode, "Abs", "%T" or "E" (Fig 43).





 Put the blank cuvette into the Reference Light Path and Main Light Path, press 【ZERO】 to scan the base line (Fig 44). Press the key 【ESC/STOP】 to stop scanning;

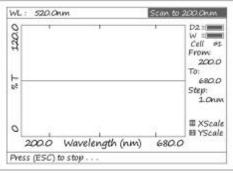
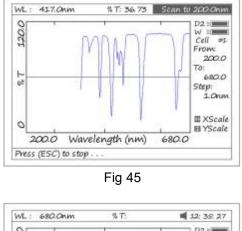
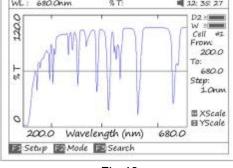


Fig 44

4. Put the sample cuvette into Main Light Path, press **[START]** to scan the sample(Fig 45) **[ESC/STOP]** to stop scanning. When scan has finished the beeper beeps 3 times (Fig 46).







5. If you want to change the scale, press 【<】 or 【>】 to change "x" scale (Fig 47), input upper limit and lower limit by pressing the numeric keypad. To change "y" scale press 【∧】 or 【∨】.
After these inputs the instrument will redraw the curve (Fig 48).

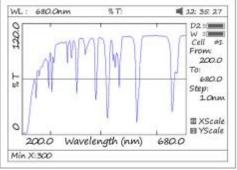
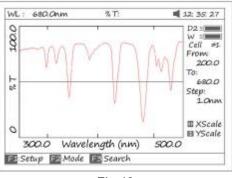
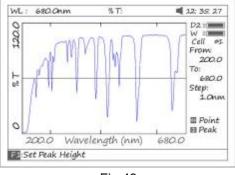


Fig47



- Fig 48
- 6. Press **[F3]** to search the Abs/%T value of the scan. There are two ways for you to search (Fig 49).





 Peak to peak, press [F1] to set "peak height" and input value by pressing the numeric keypad (Fig 50). Press [/] to search the peak from left to right and press [/] to search from right to left. The value of every peak found will be displayed on the screen one at a time (Fig 51).

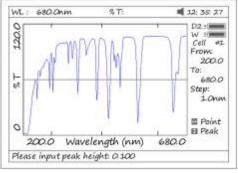


Fig 50

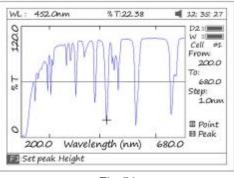


Fig 51

2) Point to point, Press [>] to search the point from left to right and press [<) to search from right to left. The search step interval is the same as the scan step. The value of every point searched will be displayed on the screen.

Save Curve

Press **[SAVE]** to save the curve. Note: Load/Save requires the first scan display page (Fig. 48). Press **ESC** if in Search to return to the required page

Print Test Report

Press **[PRINT]** to print the curve you have loaded or scanned (Fig 52).

Note: The report always is printed in Fig 46

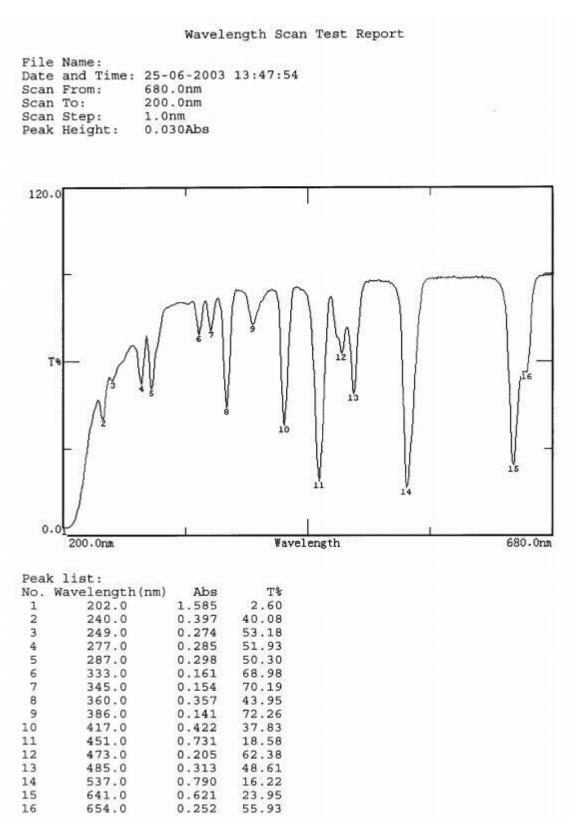
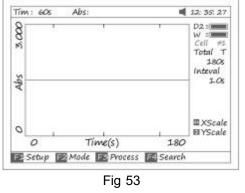


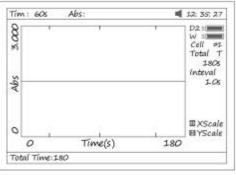
Fig 52

Kinetics

Press **[4]** in main menu for "Kinetics" (Fig 53). **[ESC/STOP]** to exit. To load a previous kinetics result, press **[LOAD]** and select a previously stored result (.kin)



- 😊 Test
 - 1. Press **[F1]** to set "Total Time", "Delay Time", "Time interval", and input the value by pressing the numeric keypad (Fig 54).





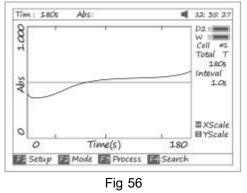
2. Select the test mode ("Abs" or "%T") by pressing [F2] (Fig 55).

3.000	,	1	-,	D2 =0 W =0 Cell #5 Total T
sąv				1805 Inteval – 1.05
0	1	Time(s)	180	III XScale III YScale



3. Set wavelength by pressing **[SET** λ **]** . Pull the blank cuvette into the Reference Light Path and Main Light Path, press **[ZERO]** for blanking

4. Pull the sample cuvette into Main Light Path, press **[START]** to scan the sample. After the delay time, the beeper beeps 3 times and time -scan starts. At the end of the time-scan, the beeper also beeps 3 times (Fig 56);



5. Press **[F3]** to process the data, and enter "Begin Time", "End Time" and "Factor" (Fig 57) and the value in I.U. will be calculated and displayed (Fig 58). The average straight line between the Begin Time and End Time will be calculated. The gradient of this line gives the rate of change of $\Delta A/min$.

Note: I.U.=Factor $\times \Delta A/min$

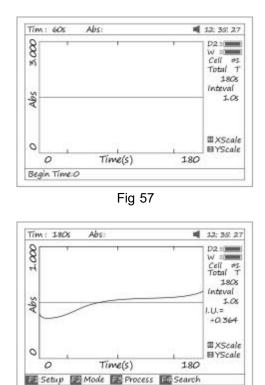


Fig 58

- 6. If you want to change the scale, please refer to step 5 of "WL scan".
- Press [F4] to search the Abs/%T value in relation to the time axis. Search point to point by pressing the key [<] or [>]. Please refer to step 6 of "WL scan".

Save Curve

Press the key **[SAVE]** to save curve. Note: Load/Save requires the first kinetics display page Fig. 56. Press **ESC** if in Search to return to the required page.

Print Test Report

Press the key [PRINT] to print the curve you have loaded or scanned (Fig 59).

Kinetics Test Report File Name: Q1.kin Date and Time: 26-06-2003 08:20:11 Total Time: 180s Time Interval: 1.0s I.U.: +0.000 From 0s to 1s

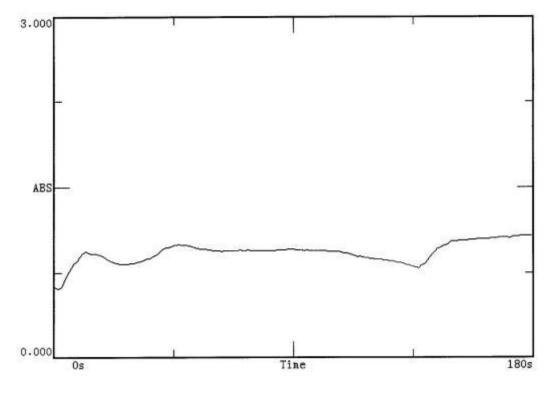
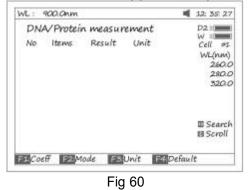


Fig 59

DNA/Protein

Press **[5]** in main menu for "DNA/Protein" (Fig 60). **[ESC/STOP]** to exit. Note:The algorithm of the test refer to Appendix A please.



To load previous DNA results, press (LOAD) and select a previously stored result (.dna)

- Contract Contract
 - 1. To use a simpler or different algorithm, you can enter your own values for f1-f4. Press **[F1]** to set f1-f4. Input the value by pressing the numeric keypad (Fig 61).

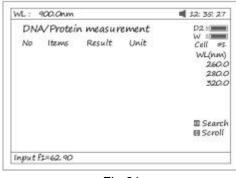


Fig 61

 Press [F2] to select test mode. "Absorbance difference 1" is for testing at the wavelength 260nm,280nm and 320nm (optional),and the "Absorbance difference 2" is for testing at the wavelength 260nm,280nm and 320nm (optional,Fig 62). Then select with/without reference. If selected with reference (no), the A ref. will be "0" (Fig 63).

ML:	900.0nm			12: 35: 27
DNA	VProtei	n measur	ement	D2 =
No	Itews	Result	Unit	W =0 Cell #1 WL(nm) 260.0 280.0
				© Search ⊡ Scrolí

Fig 62

DN	A/Protei	n measur	ement	D2 =
No	ltems	Result	Unit	W =1 Cell #1 WL(nm) 260.0 280.0
				© Search ⊡ Scrolí

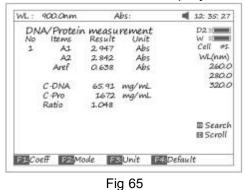
Fig 63

3. Press **[F3]** to select the unit of concentration (Fig 64).

VL :	900.0nm			12: 35: 27
DN	↓Protei	n measur	ement	D2=
No	ltews	Result	Unit	W =0 Cell #5 WL(nm) 260.0 280.0
				© Search ⊡ Scrolí



- 4. Push the blank cuvette into the Reference Light Path and Main Light Path , then press **[ZERO]** for blanking .
- 5. Pull the sample cuvette into Main Light Path, press **[START]** to test the sample. The test result will be displayed on the screen (Fig 65).



6. If there is more than one sample, repeat step 5 for the next sample.

Press the key 【<) or 【>】 for searching. Input the sample number (Fig 66), the result will be displayed on the screen. Press the key 【∧】 or 【∨】 to browse the test results one by one.

ML :	900.0nm	1	ws:	12: 35: 27
DA	IN Protein	measu	rement	D2:
No			Unit	W =0
1	AL	2.947	Abs	Cell #1
	A2.	2.842	Abs	WL(nm)
	Aref	0.638	Abs	260.0
	0.000			280.0
	C-DNA	65.92	ma/mL	320.0
	C-Pro	1672	mg/mL	
	Ratio	1.048		
				I Search
				El Scrolí
Searc	h sample:3			

Fig 66

Recall the default

Press the key [F4] to recall the default of the f1-f4.

Save Data

Press the key **[SAVE]** to save data.

Print Test Report

Press the key **[PRINT]** to print the test result (Fig 67).

DNA / Protein Test Report

File Name: Date and Time: 26-06-2003 09:16:33

No	260.0nm	280.0nm	320.0nm	C-DNA	C-Pro	Ratio
1	0.226	0.212	0.102	3.825	76.60	1.127
2	0.226	0.213	0.102	3.803	79.32	1.113

Unit:ug/mL

Fig 67

Multi Wavelength

Press [6] in main menu for "Multi WL" (Fig 68). [ESC/STOP] to exit.

WL:	900.0nm	Abs:	12: 35: 27
Mu	lti Waveleng	th Test	D2 =
No	WL(nm) <i>500.0</i>	Abs	Cell #1 I WL
			III Search III Scroli
FEIW	setup 📴	10de	
		Fig 68	

To load previous Multi Wavelength results, press **(LOAD)** and select previously stored results (.mul)

- 😊 Test
 - Press **[F1]** to setup a group of wavelengths for testing by pressing the numeric keypad followed by **[ENTER]**. (^) or (^Y) to modify the inputted data Fig. 69. Press **[ESC/STOP]** to finish setup and exit. Note: It is recommended to enter the highest wavelength first.

WL :	900.0nm	Abs:	12: 35: 27
Mi No	11ti Waveleng WL(nm) 500.0 400.0	th Test Abs	D2.5 W 3 Cell #1 2 WL
			≣ Search ≣ Scroll



2. Press **[F2]** to select mode (Fig 70).

WL: 900.0nm	Abs:	12: 35: 27
Multi Waveleng No WL(nm) 500.0 400.0 300.0	gth Test Abs	D2 = W = Call = 3 WL
		⊠ Search ₪ Scroli
Please select mode:	Abs	
	Fiq70	

3. Push the blank cuvette into the Reference Light Path and Main Light Path, then press 【ZERO】 for Blanking .

4. Pull the sample cuvette into Main Light Path, press **[START]** to test. The test results will be displayed on the screen (Fig 71).

500.0nm	Abs:	12: 35: 27
Iti Wavelend	th Test	D2 ≊
	Abs	W B
500.0	0.87	Cell #1 3 WL
400.0	0.42	5 WL
300.0	0.81	
		₪ Search ₪ Scroll
	WL(nm) 500.0 400.0	(Iti Wavelength Test WL(nm) Abs 500.0 0.87 400.0 0.42

Fig 71

- 5. If there is more than one sample, repeat step 4 for the next sample. **Note:** When the test has finished, the wavelength will go to the first WL.
- Press 【<) or 【>】 for searching. Input the sample number, the result will be displayed on the screen. Press 【∧】 or 【∨】 to browse the test results one by one.

Save Data

Press **[SAVE]** to save data.

Print Test Report

Press [PRINT] to print the test results (Fig 72).

Multi-Wavelength Test Report

File Name: M1.mul Date and Time: 26-06-2003 09:25:16

No	300.0nm	400.0nm	500.0nm
1	0.107	0.074	0.054
2	0.106	0.073	0.055
3	0.106	0.072	0.054

Unit:Abs

Fig 72

Setting and Calibration

Utility

Press [7] in Main menu for "Utility" (Fig 73). [ESC/STOP] to exit.

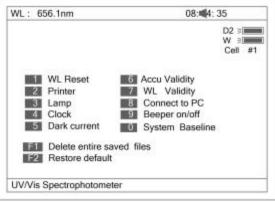


Fig 73

WL Reset

Press [1] to reset wavelength (Fig74).

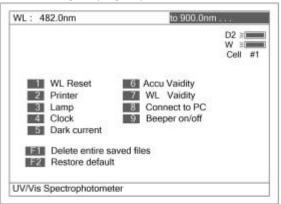


Fig 74

PrinterPress [2] to set printer (Fig 75). [ESC/STOP] to exit.

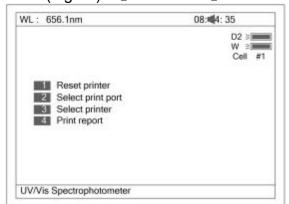
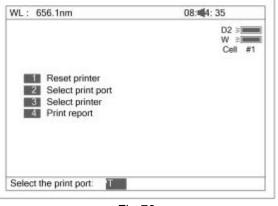


Fig 75

- 1. Press **[1]** in Fig 75 to Reset Printer.
- 2. Press **[2]** in Fig 75 to select print port (LPT or Comm., Fig 76).





3. Press **[3]** in Fig 75 to select printer (HP PCL (1 colour cartridge), PCL (black mode), Epson ESC/P or Epson/P2 or above, Fig77).

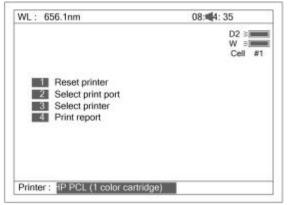


Fig 77

4. Press **[4]** in Fig 75 to select print mode. If you select "Print screen" mode, a little icon will be displayed on the top line of the screen (Fig 78), if you select "Print report" mode, the little icon will disappear.

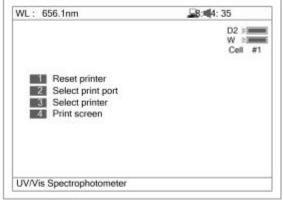
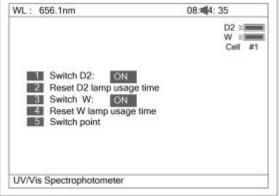


Fig78







1. Press [1] in Fig 79 to switch on/off D2. Fig 80.

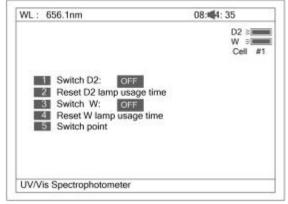


Fig 80

2. Press **[2]** in Fig 79 to reset usage time of D2(Fig 81). Press **[**∧**]** or **[**∨**]** to select "Yes" or "No", and then press **[ENTER]**.

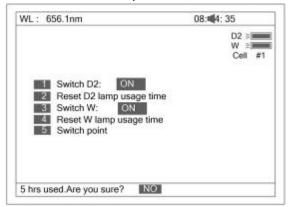


Fig 81

3. Press **[3]** in Fig 79 to switch on/off W. The indication is also on the top right corner of the screen (Fig 82).

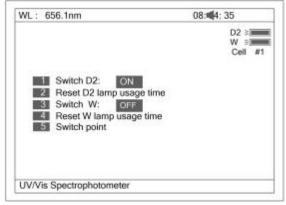


Fig 82

4. Press **【4】in Fig 79** to reset usage of W (Fig 83). Press **【**∧**】** or **【**∨**】** to select "Yes" or "No", and then press **【ENTER】**.

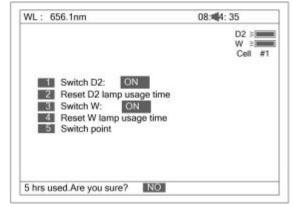


Fig 83

5. Press **[5]** in Fig 79 to set the switch usage point of D2 and W lamp (Fig 84).

D2 ∋ W ∋ Cell #1

Fig 84

Clock

Press **[4]** In Fig73 to set the display mode and modify the clock (Fig 85). **[ESC/STOP]** to exit.

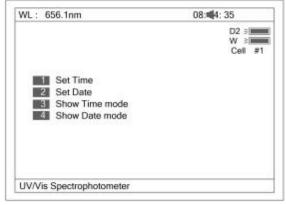


Fig 85

1. Press **[1]** in Fig 85 to modify time by pressing the numeric keypad (Fig 86).

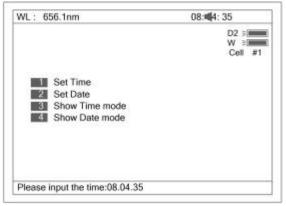


Fig 86

- 2. Press **[2]** in Fig 85 to modify date by pressing the numeric keypad.
- 3. Press **[3]** in Fig 85 to set the date display on the top right corner of the screen.

4. Press **[4]** in Fig 85 to set the time display on the top right corner of the screen (Fig 87).

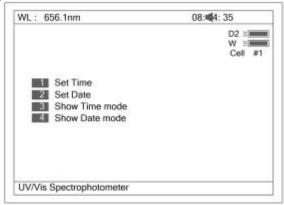


Fig 87

Dark Current

Press [5] In Fig73 to get dark current (Fig 88).

VL: 656.1nm	ark current
	D2 ≓ W ⊨ Cell #1
WL Reset	Accu Validity
2 Printer	WL Validity
Lamp	Connect to PC
Clock	Beeper on/off
5 Dark current	
Delete entire	saved files
Restore defa	tuit

Fig 88

Accu Validity
 Press 【4】 In Fig73 to do accu validity (Fig 89). 【ESC/STOP】 to exit.

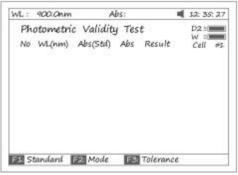


Fig 89

1. Press **[SET** λ] to set the wavelength. Press **[ENTER]** to edit and input wavelength by pressing the numeric keypad (Fig 90). **[ESC/STOP]** to finish inputting and exit.

WL :	900.0nm	Ab	\$1	12:35	27
Ph No I 2 3	otometric WL(nm) 440.0 9600 more-	Validity Abs(Std)	Test Abs Resul	D2 ±≣ t W ≊ Cell	_
Pleas	e input wl:	546			

2. Press **[F1]** to set the standard value, Press **[ENTER]** to edit and input by pressing the numeric keypad (Fig 91). **[ESC/STOP]** to finish inputting and exit.

WL :	900.0nm	%7	ē.	12:35	6 27
No 1	otometric WL(nm) 440.0 546.0 635.0	% T(Std) 9.28		D2 ≥i W ≊i Cell	
				BS	elect
Input	the standa	nd:0			

Fig 91

3. Press **[F2]** to select test mode (Abs or %T, Fig 92).

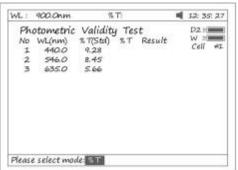


Fig 92

4. Press **[F3]** to set tolerance (Fig 93).Input the value by pressing the numeric keypad.

WL :	900.0nm	%7	9/1	12: 35: 27
Ph No 1 2 3	otometric WL(nm) 500.0 500.0 700.0	% T(Std) 9.28 8.45	Test %T Result	D2:5 ₩ 30 Cell +1
Input	tolerance c	0.005		



- 5. Press **[ZERO]** for Blanking.
- 6. Put the sample (calibrated neutral density filter) into Main Light Path. Press **[START]** to check. The results will be displayed on the screen (Fig 94). If the discrepancy between the results and the calibrated standards is not more than the tolerance, "pass" will be displayed after the test result. Otherwise, "fail" will be displayed.
- 7. The result can be saved, loaded and printed by pressing [SAVE] , [LOAD] and [PRINT].

WL:	900.0nm	9	T:		12:35	27
	otometric WL(nm) 500.0 600.0 700.0	% T(Std) 9.28 8.45	%T 9.28	Result pass	D2 =0 W =0 Cell	-
F1 S	tandard 👖	2 Mode	F3	Toleranci	ı	

Fig 94

WL Validity

Press **[7] in Fig 73** to WL validity (Fig 95). **[ESC/STOP]** to exit.

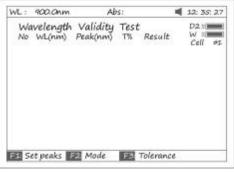


Fig 95

Press **[F1]** to set the standard peak. Press **[ENTER]** to edit and input wavelength by pressing the numeric keypad (Fig96). **[ESC/STOP]** to finish inputting and exit.

WL :	900.0nm	27	r:	-	12: 35: 2
		Validity Ptak(nm)		Result	D2 = W = Cell ≠
ę	aU 1.4				🖩 Selei



2. Press **[F2]** to select test mode (Abs or %T, Fig 97).

WL :	900.0nm	%	T:	4	12: 35: 21
No	Welength WL(nm) 2414 3610 417.0 537.6 6414 807.4	Validity Peak(nm)	Test %T	Result	D2:0 W 0 Call #3
Pleas	e select mo	de: %T			

3. Press **[F3]** to set tolerance (Fig 98). Input the value by pressing the numeric keypad.

WL :	900.0nm	27	r:		12: 35: 27
N0 1 2 3 4 5	Welength WL(nm) 241.4 361.0 417.0 537.6 641.4		Test %T	Result	D2:: W : Cell #1-
6	807.4				🗈 Select

Fig 98

4. Press **[ZERO]** for blanking.

5. Put the sample (calibrated holmium liquid) into Main Light Path. Press **[START]** to check. The results will be displayed on the screen (Fig 99). If the discrepancy between the results and the calibrated values is not more than the tolerance, "pass" will be displayed after the test results. Otherwise, "fail" will be displayed.

WL: 4	100.0nm	%T:		ي 🕨 🗧	12: 35	27
Wo	velength	Validity	Test		D2:0	=
No	WL(nm)	Peak(nm)	2 T	Result	Cell	61
1		239.7			L CH	
		360.4	42.82	pass		
23	417.0	416.9	37.63	pass		
4	537.6 641.4	537.2	16.5D	pass		
5	641.4	641.3	24.33	pass		
6	807.4	807.7	92.38	pass		
1 Sta	ndard 😰	Mode	Tolen	ance		_
		Fig	00			

The result can be saved, loaded and printde by pressing 【SAVE】,
 【LOAD】 and 【PRINT】

Connect to PC

Press [8] in Fig 73 to connect to PC (Fig 100). If the instrument is controlled by PC, the screen displays as Fig 100A. Press **[ESC/STOP]** to exit.

WL: 656.1nm	_18:44: 35
	D2 ∋t W ∋t
Connecting to computer	
UV/Vis Spectrophotometer	



WL: 656.1nm	_18:=1 4: 35
	D2 ≶ ₩ 8
Controlled by PC	
UV/Vis Spectrophotometer	

Fig 100A

Beeper on/off

Press [9] in Fig 73 to turn on/off the beeper

Delete entire saved files

Press **[F1]** in Fig 73 to delete entire saved files. After the delete the files, double confirm need to do.

Restore default

Press **[F2]** in Fig 73 to restore the default parameters.

Appendix A

	DNA/Pr	otein I	est Algorithi	n	
Test Name	Method	Wavelength(s)	Calculations	Parameters	Displayed Units
DNA MEASU	UREMENT				
DNA/Protein Concentration and DNA purity	Absorbance difference (260,280) Absorbance difference (260,230)	A1=A260nm A2=A280nm Aref=A320nm (optional) A1=A260nm A2=A230nm Aref=A320nm (optional)	DNA concentration: $(A_1-A_{ref})f_1-(A_2-A_{ref})f_2$ Protein concentration $(A_2-A_{ref})f_3-(A_1-A_{ref})f_4$ DNA concentration: $(A_1-A_{ref})f_1-(A_2-A_{ref})f_2$ Protein concentration $(A_2-A_{ref})f_3-(A_1-A_{ref})f_4$	$\begin{array}{c} f_1 = 62.9 \\ f_2 = 36.0 \\ f_3 = 1552 \\ f_4 = 757.3 \end{array}$ $f_1 = 49.1 \\ f_2 = 3.48 \\ f_3 = 183 \\ f_4 = 75.8 \end{array}$	DNA: µg/ml Protein: µ g/ml
	Absorbance ratio	A1=A260nm A2=A280nm or A230nm Aref=A320nm (optional)	$Ratio = \frac{A_1 - A_{ref}}{A_2 - A_{ref}}$	None	No units(ratio)

DNA/Protein Test Algorithm

Appendix B

A number of correction techniques can be used to eliminate or reduce interference errors. In general, if the source of the error is known and is consistent from sample to sample, the error can be eliminated. On the other hand, if the source is unknown and varies from sample to sample, the error can be reduced but not eliminated. Correction techniques can always require data from at least two wavelengths. The more sophisticated correction techniques require multiwavelength or spectral data.

A.1 Isoabsorbance

When a known interfering component with a known spectrum is present, the error introduced by this component at the analytical wavelength for the target analyte can be eliminated by selecting a reference wavelength at which the interfering compound exhibits the same absorbance as it does at the analytical wavelength. The absorbance at this reference wavelength is subtracted from the absorbance at the analytical wavelength, as shown in Figure A1.The residual absorbance is the true absorbance of the analyte.

This technique is less reliable when the spectra of the analyte and of the interferent are highly similar. Moreover, it can correct for only one interference

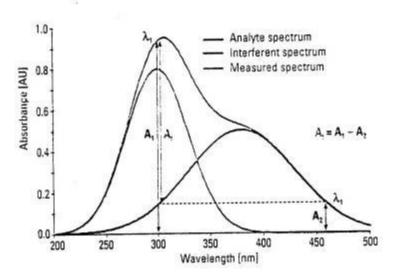


Fig A1 Isoabsorbance correction

A.2 Three-point correction

The three-point, or Morton-Stubbs correction uses two reference wavelengths, usually those on either side of the analytical wavelength. The background interfering absorbance at the analytical wavelength is then estimated using linear interpolation (see Figure A2).This method represents an improvement over the single-wavelength reference technique because it corrects for any background absorbance that exhibits a linear relationship to the wavelength. In many cases, if the wavelength range is narrow, it will be a reasonable correction for non-linear background absorbances such as that resulting from scattering of from a complex matrix.

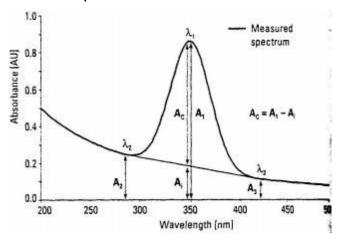


Fig A2