

HEWLETT-PACKARD

HP-67/HP-97

Clinical Lab and Nuclear Medicine Pac



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WE NEED YOUR HELP

To provide better calculator support for people like you, we need your help. Your timely inputs will enable us to provide high quality software in the future and improve the existing application pacs for your calculator. Your early reply will be extremely helpful in this effort.

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2. How important was the availability of this pac in making your decision to buy a Hewlett-Packard calculator? ☐ Would not buy without it. ☐ Important
☐ Not important
3. Did you buy this pac and your calculator at the same time? ☐ Yes ☐ No
4. In deciding to buy this application pac, which three programs seemed most useful to you? Program numbers 1. _____ 2. _____ 3. _____
5. Which three programs in this application pac seemed least useful to you?
Program numbers 1. _____ 2. _____ 3. _____
6. What program(s) would you add to this pac?

7. In the list below, select up to three application areas for which you purchased this pac. Please indicate the order of importance by 1, 2, 3, (1 represents the most important area).

- | | |
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| <input type="checkbox"/> 32 Chemistry | Other |
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| <input type="checkbox"/> 37 Statistics | <input type="checkbox"/> 75 Navigation |
| <input type="checkbox"/> 39 Other (Specify) _____ | <input type="checkbox"/> 79 Other (Specify) _____ |

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ATTENTION: APPLICATIONS

Introduction

The 19 programs of Clinical Lab and Nuclear Medicine Pac have been drawn from the fields of clinical chemistry, nuclear medicine, radioimmunoassay, and statistics.

Each program in the pac is represented by a magnetic program card and a section in this manual. The manual provides a description of the program with relevant equations, a set of instructions for using the program, and one or more example problems, each of which includes a list of the actual keystrokes required for its solution. Program listings for all the programs in the pac appear at the back of this manual. Explanatory comments have been incorporated in the listings to assist you should you want to study the actual workings of the program.

No knowledge of programming is required to use the programs in this pac. However, some familiarity with keyboard operations, as described in Sections 1 through 5 of the Owner's Handbook, is assumed. If you have already run a number of programs from Standard Pac or another applications pac, you will be able to use these programs with very little additional instruction. We recommend that you read only "A Word about these Programs" on pages iv and v of this manual. If, on the other hand, this is your first exposure to running pre-recorded programs, be sure to read the entire introductory section on pages iv to xii.

We hope that Clinical Lab and Nuclear Medicine Pac will assist you in the solution of numerous problems around the laboratory. We have tried to provide you with the most commonly used statistics programs as well, but should you find the need for more, there is another pac, Stat Pac I, exclusively for statistics.

We would very much appreciate knowing your reactions to the programs in this pac, and to this end we have provided a questionnaire inside the front cover of this manual. Would you please take a few minutes to give us your comments on these programs? It is in the comments we receive from you that we learn how best to increase the usefulness of programs like these.

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| Program | Page |
|--|-------------|
| Clinical Chemistry | |
| 1. Beer's Law | .01-01 |
| Converts between absorbance and % transmittance; solves for an unknown concentration given a standard concentration and the absorbance or % transmittance of the standard and unknown. | |
| 2. Protein Electrophoresis | .02-01 |
| Given integration counts of a number of protein fractions, finds percentage of each. Calculation of weights optional. | |
| 3. LDH Isoenzymes | .03-01 |
| Given values for the five LDH isoenzymes, finds activity of each as a percent of total. Compares results against normal values. | |
| 4. Body Surface Area | .04-01 |
| Calculates an estimated BSA by method of Dubois or Boyd. Accepts either English or metric units. | |
| 5. Urea Clearance | .05-01 |
| Calculates urea clearance with option of correcting for BSA. | |
| 6. Creatinine Clearance | .06-01 |
| Calculates creatinine clearance with option of correcting for BSA. | |
| 7. Amniotic Fluid Assay | .07-01 |
| Performs calculations for the spectrophotometric estimation of bile pigments in amniotic fluid. | |
| 8. Blood Acid-Base Status | .08-01 |
| Finds total plasma CO ₂ and base excess from PCO ₂ , pH and Hgb concentration. | |
| 9. Oxygen Saturation and Content | .09-01 |
| Finds oxygen saturation and content in blood given PO ₂ , PCO ₂ , pH, and body temperature. | |
| 10. Red Cell Indices | .10-01 |
| Given hematocrit percent, red cell count, and hemoglobin, finds mean corpuscular volume, mean corpuscular hemoglobin, and mean corpuscular hemoglobin concentration. | |
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| 12. Schilling Test | .12-01 |
| The radioisotope determination of vitamin B ₁₂ absorption. | |
| 13. Thyroid Uptake | .13-01 |
| The radioisotope determination of thyroid uptake. | |
| 14. Radioactive Decay Corrections | .14-01 |
| Finds the activity of a radioisotope corrected for decay over time. | |
| Radioimmunoassay | |
| 15. Radioimmunoassay | .15-01 |
| Computes least-squares regression line of logit of net counts vs. log concentration, including regression constants, correlation coefficient, and concentration for a given count. | |

Statistics

- 16. **Basic Statistics****16-01**
Computes mean, standard deviation, standard error, and coefficient of variation for grouped or ungrouped data.
- 17. **Chi-square Evaluation and Distribution****17-01**
Computes the chi-square statistic for goodness of fit. For given $x \geq 0$, finds the chi-square density function $f(x)$ and the cumulative distribution $P(x)$.
- 18. **t Statistics****18-01**
Computes the paired t statistic and the unpaired t statistic.
- 19. **t Distribution****19-01**
For a given $x > 0$, evaluates the t density function and cumulative distribution.

A WORD ABOUT THESE PROGRAMS

This application pac has been designed for both the HP-97 Programmable Printing Calculator and the HP-67 Programmable Pocket Calculator. The most significant difference between the HP-67 and the HP-97 calculators is the printing capability of the HP-97. Most of the computed results in this pac are output by the command PRINTx. On the HP-97 these results will be output on the printer. On the HP-67 each PRINTx command will be interpreted as a PAUSE: the program will halt, display the result for about five seconds, then continue execution.

If you use an HP-67, you may want more time to copy down the number displayed by a PRINTx command. All you need to do is press any key on the keyboard during the pause interval in which the result is displayed. This action will cause the program to halt; execution of the halted program may be re-initiated by pressing **R/S**. Values that are output by a PRINTx command are marked by three asterisks (***) in the keystroke solutions to example problems. The keystroke solutions reflect another slight difference between the HP-67 and the HP-97. It is sometimes necessary in these solutions to include operations that involve prefix keys, namely, **f** on the HP-97 and **f**, **g**, and **h** on the HP-67. For example, the operation 10^x is performed on the HP-97 as **f** 10^x and on the HP-67 as **g** 10^x . In such cases, the keystroke solution omits the prefix key and indicates only the operation (as here, 10^x). As you work through the example problems, take care to press the appropriate prefix keys (if any) for your calculator.

Programs 1 through 13 of this pac are alike in that many of the same operations are available in each of these programs. A look at the magnetic cards for these programs will show three instructions repeated in gold on every card of these thirteen: PTNT #, P OFF?, and REPRINT. These three operations are intended primarily for use on the HP-97. In addition, either CLEAR or START appears on all of the first thirteen cards. Some discussion of these common operations may be helpful.

The instruction PTNT # allows you to key in a patient number which will be immediately printed in order to identify the data and results of the following calculations. The patient number used should be a whole number; the program will append two digits after the decimal point to identify the program being used, 01 to 13. For example, if the patient number 1234 is used in program 7, the program would print the identification 1234.07, which serves to identify the entire context of the calculations which are to follow. The use of the patient number for identification purposes is entirely optional and may be omitted.

The interrogative P OFF? asks the question: do you want to turn the print function off? When the program is loaded, a flag is set that causes all inputs and outputs of the program to be printed. If this information is not all desired, you may eliminate some or all of it, depending on the program, by turning the print

function off. It may later be turned back on at any time without affecting the operation of the program.

The instruction REPRINT allows for an additional printout of all data and results after a calculation has been completed. Frequently in the clinical lab, the results of a test must be reported to several different departments. The REPRINT feature allows you to obtain additional copies of the data and results directly from the program.

The instructions CLEAR and START are similar in that both have to do with initialization of the program and should be executed before any other operation in the program. They differ in that CLEAR is an optional instruction and START is mandatory. Basically, CLEAR simply sets certain registers to zero to insure that meaningless information is not output during a REPRINT. On the other hand, START loads registers with necessary initial values without which the program would fail to function properly.

RUNNING A PROGRAM

Loading a Program

Select the *Protein Electrophoresis* card, CL1-02A, from the card case supplied with this application pac.

Set the PRGM-RUN switch to RUN.

If you are using the HP-97, set the printer switch to MAN. All the programs in this pac are designed for manual printer setting.

Gently insert either end of the card (printed side up) in the reader slot of your calculator as shown in figure 1a or 1b.



Figure 1a. HP-97

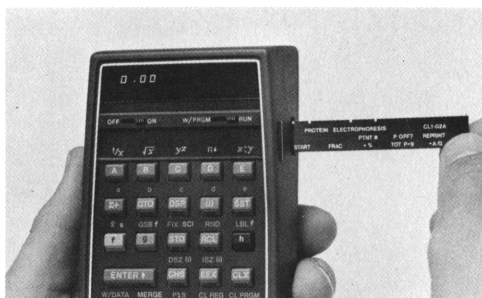


Figure 1b. HP-67

When the card is part way in, a motor engages and passes it out the other side of the calculator. Sometimes the motor engages but does not pull the card in. If this happens, push the card a little farther into the machine. Do not impede or force the card; let it move freely.

The display will show “Error” if the card reads improperly. In this case, press **CLX** and reinsert the card.

Since *Protein Electrophoresis* is longer than 112 steps (the capacity of one side of a magnetic card), the display now shows “Crd” indicating that a

second card pass is necessary to load the remaining steps. With the writing still visible to you, insert the *opposite* end of the card (figures 2a and 2b) and pass the card through the card reader again.

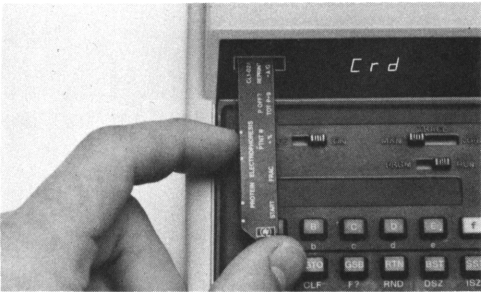


Figure 2a. HP-97



Figure 2b. HP-67

When the motor stops, remove the card from the other side of the calculator and insert it in the “window slot” of the calculator (figures 3a and 3b).



Figure 3a. HP-97

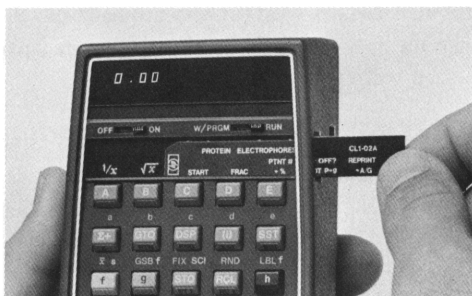


Figure 3b. HP-67

The program has now been stored in the calculator. It will remain stored until another program is loaded or the calculator is turned off.

The Magnetic Card

Complete instructions for running the program are found in the User Instructions form for that program. The first few times you run the program, you should refer to these instructions at each step of the operation. Thereafter, mnemonic symbols on the magnetic card itself will provide shorthand instructions to the program's operation.

Take a look at the card that you have inserted in the window slot of the calculator. Notice that the mnemonic symbols on the card are grouped above the user-definable keys **A** through **E**. For example, the symbols “ $\rightarrow\%$ ” and “PTNT #” are associated with key **C**. Symbols in gold are associated with the shifted keys **f A** through **f E**.

Below is a table of the important symbols and conventions you will find on magnetic cards.

| SYMBOL OR CONVENTION | INDICATED MEANING |
|--|---|
| White mnemonic: x A | White mnemonics are associated with the user-definable key they are above when the card is inserted in the calculator's window slot. In this case the value of x could be input by keying it in and pressing A . |
| Gold mnemonic: y x f E x \uparrow y A | Gold mnemonics are similar to white mnemonics except that the gold f key must be pressed before the user-definable key. In this case y could be input by pressing f E . \uparrow is the symbol for ENTER\uparrow . In this case ENTER\uparrow is used to separate the input variables x and y. To input both x and y you would key in x, press ENTER\uparrow , key in y and press A . |

SYMBOLS AND CONVENTIONS (Continued)

| SYMBOL OR CONVENTION | INDICATED MEANING |
|-----------------------------------|---|
| \boxed{x} A | <p>The box around the variable x indicates input by pressing STO A.</p> |
| (x) A | <p>Parentheses indicate an option. In this case, x is not a required input but could be input in special cases.</p> |
| $\rightarrow x$ A | <p>\rightarrow is the symbol for calculate. This indicates that you may calculate x by pressing key A.</p> |
| $\rightarrow x, y, z$ A | <p>This indicates that x, y, and z are calculated by pressing A once. The values would be printed in x, y, z order.</p> |
| $\rightarrow x; y; z$ A | <p>The semi-colons indicate that after x has been calculated using A, y and z may be calculated by pressing R/S.</p> |
| $\rightarrow "x," y$ A | <p>The quote marks indicate that the x value will be "paused" or held in the display for one second. The pause will be followed by the display of y.</p> |
| $\diamond x$ A | <p>The two-way arrow \diamond indicates that x may be either output or input when the associated user-definable key is pressed. If numeric keys have been pressed between user-definable keys, x is stored. If numeric keys have not been pressed, the program will calculate x.</p> |
| <p>P?</p> A | <p>The question mark indicates that this is a mode setting, while the mnemonic indicates the type of mode being set. In this case a print mode is controlled. Mode settings typically have a 1.00 or 0.00 indicator displayed after they are executed. If 1.00 is displayed, the mode is on. If 0.00 is displayed, it is off.</p> |
| <p>START</p> A | <p>The word START is an example of a command. The start function should be performed to begin or start a program. It is included when initialization is necessary.</p> |
| <p>DEL</p> A | <p>This special command indicates that the last value or set of values input may be deleted by pressing A.</p> |

FORMAT OF USER INSTRUCTIONS

The completed User Instructions Form—which accompanies each program—is your guide to operating the programs in this Pac.

The form is composed of five labeled columns. Reading from left to right, the first column, labeled STEP, gives the instruction step number.

The INSTRUCTIONS column gives instructions and comments concerning the operations to be performed.

The INPUT DATA/UNITS column specifies the input data, and the units of data if applicable. Data input keys consist of **[0]** to **[9]** and decimal point (the numeric keys), **[EEX]** (enter exponent), and **[CHS]** (change sign).

The KEYS column specifies the keys to be pressed after keying in the corresponding input data.

The OUTPUT DATA/UNITS column specifies intermediate and final outputs and their units, where applicable.

The following illustrates the User Instruction Form for *Protein Electrophoresis*, CL1-02A.

| STEP | INSTRUCTIONS | INPUT DATA/UNITS | KEYS | OUTPUT DATA/UNITS |
|------|------------------------------------|--------------------|----------------|-------------------|
| 1 | Load side 1 and side 2. | | | |
| 2 | Initialize. | | [A] | 0.00 |
| 3 | (optional) Key in patient number. | Ptnt # | [f] [C] | Ptnt # .02 |
| 4 | To suppress output of data, turn | | | |
| | print function off. | | [f] [D] | 0.00 |
| 5 | To turn print back on later. | | [f] [D] | 1.00 |
| 6 | Key in the counts of the first | | | |
| | protein fraction. | Fract ₁ | [B] | 1.00 |
| 7 | Repeat this step for the rest of | | | |
| | the fractions. | Fract _i | [B] | i |
| 8 | Calculate the percentage each | | | |
| | fraction is of the whole. | | [C] | % |
| 9 | (optional) Key in the total grams | | | |
| | of protein and find the grams in | | | |
| | each fraction. | Total Protein | [D] | grams |
| 10 | (optional) Find the albumin/ | | | |
| | globulin ratio. | | [E] | A/G |
| 11 | (optional) Obtain a reprint of all | | | |

| STEP | INSTRUCTIONS | INPUT DATA/UNITS | KEYS | OUTPUT DATA/UNITS |
|------|---------------------------------|------------------|-------------------|------------------------|
| | data and results (Total Protein | | | |
| | and grams omitted if Total | | | |
| | Protein not input). | | f E | Ptnt # .02 |
| | | | | Fract ₁ ... |
| | | | | Fract _n |
| | | | | (%) ₁ ... |
| | | | | (%) _n |
| | | | | Total Protein |
| | | | | grams ₁ ... |
| | | | | grams _n |
| | | | | A/G |
| 12 | For a new case, go to step 2. | | | |

Since you loaded this program in “Loading a Program” on page vi, step 1 is already done and we can move to step 2. (If you turned your calculator off, you must reload the program.) Leave the magnetic card in the window slot above keys **A** through **E**.

Step 2 is an initialization procedure, marked START on the magnetic card. Press **A** now to perform the initialization, as shown in the KEYS column. You should see a display of 0.00.

Step 3 is optional and allows for input of the patient number if identification of the output is desired. The number output at this step is the patient number followed by “.02”, which marks the second program of the pac, CL1-02A. Key in the patient number 1234 and see an output of 1234.02.

Steps 4 and 5 have to do with the optional print mode, which may be turned off or on through the keystrokes **f** **D**. When the program is loaded, the print function is on; pressing **f** **D** will turn it off and display 0.00. Try it. Successive presses of **f** **D** will turn the print function on, then off, alternately displaying 1.00 (on) and 0.00 (off). Try this, but leave 1.00 displayed (print function on) when you are finished. This will allow the input data to be output through PRINTx commands.

Step 6 begins the actual input of the fractionation data. You are to key in the counts for the first protein fraction (Fract₁ under INPUT DATA/UNITS) and press **B**. This value will be output and a 1.00 will be displayed to mark the input of the first fraction. Step 7 instructs you to input the remaining protein fractionation counts in a like manner, keying in each value and pressing **B**.

The number displayed after each value is input indicates the number of functions input so far. Try this sequence with the values from the table below.

| Fraction | Substance | Counts |
|----------|----------------------|--------|
| 1 | Albumin | 67 |
| 2 | α_1 -globulin | 4 |
| 3 | α_2 -globulin | 10 |
| 4 | β -globulin | 14 |
| 5 | γ -globulin | 13 |

Use the keystrokes 67 **B** 4 **B** 10 **B** 14 **B** 13 **B**. At the end of this sequence the display should show 5.00.

Now that all fractions have been input, step 8 instructs you to find the percentages for the fractions input by pressing **C**. Each percentage is output by a PRINTx command, and the percentages will be output in the order the fractions were input. Press **C** now. The outputs you should see are, in this order, 62.04, 3.70, 9.26, 12.96, and 12.04.

Step 9 is optional. Here you may key in the total grams of protein and press **D** to find the number of grams in each fraction. Key in 7, press **D**, and you should see these outputs: 4.34, 0.26, 0.65, 0.91, and 0.84.

Step 10 is optional. You may press **E** to compute the albumin/globulin ratio. Press **E** now and find an A/G value of 1.63.

Step 11 is also optional. This is the REPRINT feature described on page v. If **f E** is pressed, the entire set of data and results will be output through PRINTx commands in the order shown in the OUTPUT DATA/UNITS column. You may do this now and check that the values returned by the REPRINT function are the same as those you keyed in or calculated earlier.

If your answers agree with ours, you are ready to try other programs in this pac. Otherwise, go back to the start of this section and try the procedure again.

Notes

BEER'S LAW



This program combines two independent routines in the area of spectrophotometry. The first routine, on keys **A** and **B**, solves Beer's law interchangeably to find either absorbance (optical density) or percent transmittance (%T). To find %T, key in absorbance and press key **A**. The output will be %T. To find absorbance, key in %T and press key **B**. Absorbance will be output.

The second routine, on keys **C**, **D**, and **E**, allows calculation of the concentration of an unknown given the concentration of a standard and the absorbance of %T of the standard and unknown. If the percent transmittance of the standard (%T_s) is known, it may be keyed in to key **C**. If the absorbance of the standard (A_s) is known instead, it may be keyed in *as a negative number* to key **C**. Similarly, for the unknown, percent transmittance (%T_u) may be keyed in as a positive number or absorbance (A_u) as a negative number to key **D**. Then the concentration of the standard (c_s) should be keyed in to key **E**. This will allow output of the concentration of the unknown (c_u).

Equations:

$$A = 2 - \log \%T$$

$$\%T = 10^{2-A}$$

$$c_u = c_s \times \frac{A_u}{A_s}$$

Reference:

Clinical Chemistry, ed. Henry, Cannon, and Winkelman, Harper and Row, 1974.

| STEP | INSTRUCTIONS | INPUT DATA/UNITS | KEYS | OUTPUT DATA/UNITS |
|------|--|------------------|-------------------|-------------------|
| 1 | Load side 1 and side 2. | | | |
| 2 | (optional) Initialize for reprint. | | F A | 0.00 |
| 3 | (optional) Key in patient number. | Ptnt # | F C | Ptnt # .01 |
| 4 | To suppress printing of data and results, turn print function off. | | F D | 0.00 |
| 5 | To turn print function back on. | | F D | 1.00 |
| 6 | To solve interchangeably for A and %T, go to step 7; to find an unknown concentration, go to step 9. | | | |
| | A\rightleftharpoons%T | | | |
| 7 | To find percent transmittance, key in absorbance. | A | A | %T |
| 8 | To find absorbance, key in percent transmittance. | %T | B | A |
| | Unknown concentration | | | |
| 9 | Key in A or %T for the standard and the unknown (follow A by CHS): | | | |
| | • Standard | $+ \%T_s(-A_s)$ | C | $+ \%T_s(-A_s)$ |
| | • Unknown | $+ \%T_u(-A_u)$ | D | $+ \%T_u(-A_u)$ |
| 10 | Key in concentration of standard and compute concentration of unknown. | C_s | E | C_u |

| STEP | INSTRUCTIONS | INPUT DATA/UNITS | KEYS | OUTPUT DATA/UNITS |
|------|-------------------------------|------------------|-------------------|-------------------------------------|
| | Reprint | | | |
| 11 | Reprint all data and results. | | f E | Ptnt # .01 |
| | | | | A |
| | | | | %T |
| | | | | +%T _s (-A _s) |
| | | | | +%T _u (-A _u) |
| | | | | c _s |
| | | | | c _u |

Example:

A standard solution with a solute concentration of 2 mg/ml is found to have an absorbance of 0.41 at 550 nm. An unknown from patient number 10183 is found to show 46% transmittance at the same wavelength. Convert this %T to absorbance. Also find the solute concentration in the unknown. After all calculations obtain a reprint.

Keystrokes:

f **A** →
10183 **f** **C** →
46 **B** →

.41 **CHS** **C** →
46 **D** →
2 **E** →

f **E** →

Outputs:

0.00 (Clear)
10183.01 *** (Ptnt ID)
46.00 *** (%T)
0.34 *** (A)
-0.41 *** (-A_s)
46.00 *** (%T_u)
2.00 *** (c_s)
1.65 *** (c_u)
10183.01 *** (Ptnt ID)
0.34 *** (A)
46.00 *** (%T)
-0.41 *** (-A_s)
46.00 *** (%T_u)
2.00 *** (c_s)
1.65 *** (c_u)

Notes

PROTEIN ELECTROPHORESIS



This program is designed to aid in the calculations of protein fractionation. The required data for the program are the integration counts for each protein fraction and, optionally, the total protein. The results calculated by the program are the percentage of the total for each fraction and, if total protein has been input, the number of grams of each protein fraction. An optional output is the albumin/globulin ratio.

To operate the program, press key **A** to initialize. Then for each fraction, key in its integration counts and press key **B**. After the counts have been keyed in for every fraction, you may press key **C** to find the percentage that each fraction is of the total. A single press of **C** will cause all the percentages to be output in the same order as the counts were input. You may then, if you wish, key in the total protein in grams, press key **D**, and output the grams of protein for each fraction.

The albumin/globulin ratio (A/G) may be calculated by pressing key **E**. If A/G is to be found, albumin should be the first fraction input, followed by the four globulin counts.

Equations:

Let Fract_i be the counts for the i^{th} fraction, and $(\%)_i$ the percentage of the total for the i^{th} fraction.

$$(\%)_i = \frac{\text{Fract}_i}{\sum_{j=1}^n \text{Fract}_j} \times 100$$

Let TPr be the total protein in grams and g_i be the number of grams of the i^{th} fraction.

$$g_i = \frac{\text{Fract}_i}{\sum_{j=1}^n \text{Fract}_j} \times \text{TPr}$$

$$\text{A/G} = \frac{\text{Fract}_1}{\sum_{j=2}^5 \text{Fract}_j}$$

Remarks:

1. If the print function is turned off, input data will not be printed. Calculated results will still be printed regardless of the status of the print function.
2. If a reprint is called for by pressing **f E**, all possible inputs and outputs will be printed except that if no value was keyed in for total protein, neither it nor the grams of each fraction will be output.
3. The use of this program need not be restricted to protein fractionation. It may be used as a general-purpose total and percent-of-total program. The only restriction is that the number of inputs (fractions) is limited to 21.

Reference:

Clinical Chemistry, ed. Henry *et. al.*, Harper and Row, 1974.

| STEP | INSTRUCTIONS | INPUT DATA/UNITS | KEYS | OUTPUT DATA/UNITS |
|------|---|--------------------|------------|-------------------|
| 1 | Load side 1 and side 2. | | | |
| 2 | Initialize. | | A | 0.00 |
| 3 | (optional) Key in patient number. | Ptnt # | f C | Ptnt # .02 |
| 4 | To suppress output of data, turn print function off. | | f D | 0.00 |
| 5 | To turn print back on later. | | f D | 1.00 |
| 6 | Key in the counts of the first protein fraction. | Fract ₁ | B | 1.00 |
| 7 | Repeat this step for the rest of the fractions. | Fract _i | B | i |
| 8 | Calculate the percentage each fraction is of the whole. | | C | % |
| 9 | (optional) Key in the total grams of protein and find the grams in each fraction. | Total Protein | D | grams |
| 10 | (optional) Find the albumin/globulin ratio. | | E | A/G |

| STEP | INSTRUCTIONS | INPUT DATA/UNITS | KEYS | OUTPUT DATA/UNITS |
|------|--------------------------------|------------------|-------------------|------------------------|
| 11 | (optional) Obtain a reprint of | | | |
| | all data and results (Total | | | |
| | Protein and grams omitted if | | | |
| | Total Protein not input). | | f E | Ptnt # .02 |
| | | | | Fract ₁ ... |
| | | | | Fract _n |
| | | | | (%) ₁ ... |
| | | | | (%) _n |
| | | | | Total Protein |
| | | | | grams ₁ ... |
| | | | | grams _n |
| | | | | A/G |
| 12 | For a new case, go to step 2. | | | |

Example:

The following integration counts are determined electrophoretically for serum proteins:

| | |
|----------------------|----|
| Albumin | 67 |
| α_1 -globulin | 4 |
| α_2 -globulin | 10 |
| β -globulin | 14 |
| γ -globulin | 13 |

If the total amount of protein is 7.0 grams, find the percentage of the total and the number of grams for each protein fraction. Also find the albumin/globulin ratio. The patient number is 10183.

Keystrokes:

| | |
|-------------------------|---|
| A | → |
| 10183 f C | → |
| 67 B | → |
| 4 B | → |
| 10 B | → |
| 14 B | → |
| 13 B | → |

Outputs:

| |
|------------------------|
| 0.00 |
| 10183.02 *** (Ptnt ID) |
| 1.00 |
| 2.00 |
| 3.00 |
| 4.00 |
| 5.00 |

C →

62.04 *** (% albumin)

3.70 *** (% α_1)

9.26 *** (% α_2)

12.96 *** (% β)

12.04 *** (% γ)

7 **D** →

7.00 *** (Total Protein)

4.34 *** (g albumin)

0.26 *** (g α_1)

0.65 *** (g α_2)

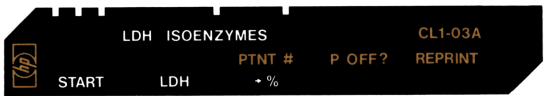
0.91 *** (g β)

0.84 *** (g γ)

E →

1.63 *** (A/G)

LDH ISOENZYMES



This program analyzes the results of the fractionation of lactic dehydrogenase isoenzymes and computes for each isoenzyme (LDH₁ through LDH₅) the percentage it represents of the whole. After key **A** is pressed to initialize the program, each enzyme value is input by keying in the value and pressing **B**. After all five LDH fractions have been input, key **C** may be pressed to find the percentage each enzyme is of the whole.

An additional feature of the program is the checking of the computed percentage of each enzyme against its accepted normal value. All five percentages are computed and output; if one or more of these values lie outside the accepted normal range, the word “Error” will be displayed at the end of all calculations. (This indicates only that a value is abnormal; the answers calculated are accurate.)

The abnormal value or values should then be determined by inspection. The normal values used by the program are shown below.

| Enzyme | Normal Range |
|------------------|--------------|
| LDH ₁ | 18%—33% |
| LDH ₂ | 28%—40% |
| LDH ₃ | 18%—30% |
| LDH ₄ | 6%—16% |
| LDH ₅ | 2%—13% |

These values for normal ranges may be changed easily within the program if you so desire. Simply look at the program listing and find the value you want to change by referring to the program comments. Delete the number as it now exists in the program and key in your own value. Do not forget to record the modified program on a blank magnetic card if you want to preserve it.

Equations:

Let LDH_i be the value of the ith LDH isoenzyme (i = 1,...,5) and LDH_i% be that enzyme’s percentage of the whole.

$$LDH_i\% = \frac{LDH_i}{\sum_{j=1}^5 LDH_j}$$

Remarks:

If the print function is turned off, input data will not be printed. Calculated results will still be printed regardless of the status of the print function.

Reference:

Clinical Chemistry, ed. Henry *et. al.*, Harper and Row, 1974.

| STEP | INSTRUCTIONS | INPUT DATA/UNITS | KEYS | OUTPUT DATA/UNITS |
|------|--|------------------|------------|--------------------|
| 1 | Load side 1 and side 2. | | | |
| 2 | Initialize. | | A | 0.00 |
| 3 | (optional) Key in patient number. | Ptnt # | F C | Ptnt # .03 |
| 4 | To suppress printing of input data, turn the print function off. | | F D | 0.00 |
| 5 | To turn the print function back on. | | F D | 1.00 |
| 6 | Key in the first LDH enzyme value. | LDH ₁ | B | 1.00 |
| 7 | Repeat step 6 for LDH values 2 through 5. | LDH _i | B | i |
| 8 | Calculate the percentage each enzyme is of the total.* | | C | LDH ₁ % |
| | | | | ... |
| | | | | LDH ₅ % |

| STEP | INSTRUCTIONS | INPUT DATA/UNITS | KEYS | OUTPUT DATA/UNITS |
|------|---------------------------------|---------------------|-------------------|----------------------|
| 9 | (optional) Obtain a reprint of | | | |
| | all data and results.* | | f E | Ptnt # .03 |
| | | | | LDH ₁ |
| | | | | ... |
| | | | | LDH ₅ |
| | | | | LDH ₁ % |
| | | | | ... |
| | | | | LDH ₅ % |
| | *A display of "Error" following | | | |
| | execution of this step | | | |
| | indicates a percentage | | | |
| | value that lies outside the | | | |
| | normal range for that | | | |
| | enzyme. | | | |

Example:

Electrophoretic separation of the LDH isoenzymes results in the following counts:

| Enzymes | Counts |
|------------------|--------|
| LDH ₁ | 95 |
| LDH ₂ | 120 |
| LDH ₃ | 85 |
| LDH ₄ | 15 |
| LDH ₅ | 22 |

Find the percentage of the whole for each isoenzyme. The patient number is 10183. Obtain a reprint of the data and results.

Keystrokes:

| | |
|----------|--------------------------|
| A | _____→ |
| 10183 | f C _____→ |
| 95 | B _____→ |
| 120 | B _____→ |
| 85 | B _____→ |
| 15 | B _____→ |
| 22 | B _____→ |

Outputs:

| |
|------------------------|
| 0.00 |
| 10183.03 *** (Ptnt ID) |
| 1.00 |
| 2.00 |
| 3.00 |
| 4.00 |
| 5.00 |

| | |
|-------------------------------|---------------------------------|
| C → | 28.19 *** (% LDH ₁) |
| | 35.61 *** (% LDH ₂) |
| | 25.22 *** (% LDH ₃) |
| | 4.45 *** (% LDH ₄) |
| | 6.53 *** (% LDH ₅) |
| | “Error” |
| CLx (clears “Error”) → | 0.00 |
| f E → | 10183.03 *** |
| | 95.00 *** |
| | 120.00 *** |
| | 85.00 *** |
| | 15.00 *** |
| | 22.00 *** |
| | 28.19 *** |
| | 35.61 *** |
| | 25.22 *** |
| | 4.45 *** |
| | 6.53 *** |
| | “Error” |

A visual scan of the results indicates that the message “Error” resulted from the percentage value of LDH₄ (4.45%) being below the normal range (6%—16%).

BODY SURFACE AREA



This program calculates body surface area by either the method of Dubois or the method of Boyd. In both cases, the required inputs are height and weight, which may be input either in metric (cm, kg) or English (in., lb.) units. Quantities in English units should be input as negative numbers; that is, **CHS** should be pressed after keying the number in.

To operate the program, the height in either cm or inches should be keyed in to **A**, and the weight in either kg or pounds keyed in to **B**. Then pressing **C** will allow the calculation of body surface area in m² by the method of Dubois; pressing **D** computes BSA in m² by the Boyd formula. Even if you have already found BSA by one method, you may also find it by the other method simply by pressing the appropriate key; the values of height and weight need not be re-input.

Equations:

Let Ht be height, Wt be weight, and BSA be the body surface area in m².

$$\text{Ht (cm)} = 2.54 \text{ Ht (in.)}$$

$$\text{Wt (kg)} = 0.45359237 \text{ Wt (lb.)}$$

Dubois:

$$\text{BSA (m}^2\text{)} = \text{Ht (cm)}^{0.725} \cdot \text{Wt (kg)}^{0.425} \cdot 7.184 \times 10^{-3}$$

Boyd:

$$\text{BSA (m}^2\text{)} = \text{Wt (g)}^{(0.7285 - 0.0188 \log \text{ Wt})} \cdot \text{Ht (cm)}^{0.3} \cdot 3.207 \times 10^{-4}$$

Remarks:

1. The Dubois formula for BSA is undefined for children with a BSA less than 0.6 m². In such cases BSA should be calculated by the Boyd formula.
2. Turning off the print function will suppress printing of both data and results.

References:

D. Du Bois and E.F. Du Bois, Clin. Cal. 10, Arch. Int. Med., 17, 863, 1916.
Edith Boyd, *Growth of the Surface Area of the Human Body*, U. of Minnesota Press, 1935, p. 132.

| STEP | INSTRUCTIONS | INPUT DATA/UNITS | KEYS | OUTPUT DATA/UNITS |
|------|---|------------------|-------------------|-------------------|
| 1 | Load side 1 and side 2. | | | |
| 2 | (optional) Initialize if reprint desired. | | f A | 0.00 |
| 3 | (optional) Key in patient number. | Ptnt # | f C | Ptnt # .04 |
| 4 | To suppress printing of data and results. | | f D | 0.00 |
| 5 | To turn print function back on. | | f D | 1.00 |
| 6 | Key in height (+ cm, - in.) | Ht | A | Ht (cm) |
| 7 | Key in weight (+ kg, - lb.) | Wt | B | Wt (kg) |
| 8 | Compute BSA by method of either | | | |
| | • Dubois | | C | BSA (m²) |
| | • Boyd | | D | BSA (m²) |
| 9 | (optional) Reprint all data and results. | | f E | Ptnt # .04 |
| | | | | Ht input |
| | | | | Wt input |
| | | | | BSA (m²) |

Example 1:

Patient number 10183 is a male, height 176 cm, weight 63.5 kg. What is his BSA by the Dubois formula? Compare by also finding the Boyd BSA.

Keystrokes:

10183 **f** **C** →
176 **A** →
63.5 **B** →
C →
D →

Outputs:

10183.04 *** (Ptnt ID)
176.00 (Ht (cm))
63.50 (Wt (cm))
1.78 *** (Dubois)
1.76 *** (Boyd)

Example 2:

Patient number 10070 is a female, height 64 inches, weight 112 pounds. Find her BSA by the Boyd formula. Obtain a reprint. Remember to input height and weight as negative numbers.

Keystrokes:

f A →
10070 f C →
64 CHS A →
112 CHS B →
D →
f E →

Outputs:

0.00
10070.04 *** (Ptnt ID)
162.56 (Ht (cm))
50.80 (Wt (kg))
1.52 *** (Boyd)
10070.04 *** (Ptnt ID)
-64.00 *** (Ht)
-112.00 *** (Wt)
1.52 *** (BSA)

Notes

UREA CLEARANCE



This program calculates urea clearance given the urine flow rate and the concentration of urea in urine and blood. The urine flow rate may be corrected for the patient's body surface area, if desired. The program will calculate standard or maximum clearance depending on whether the corrected urine flow rate is above or below 2 ml/min. The percent of mean normal may also be found.

If the urine flow rate is to be corrected for body surface area, key **f** **B** should be pressed to indicate that. No action is necessary if the correction is not desired. If correction is to be made, the program will need to find the patient's body surface area (BSA) in register R_A . If the program *Body Surface Area* (CL1-04A) has been run immediately before this program, BSA will already have been stored in R_A . Otherwise you will need to key in the patient's BSA and store it in R_A .

When inputting the urine flow rate, you may either key in the flow rate (\dot{V} , in ml/min.) directly to key **B**, or key in both the urine volume V in ml and the time t in min. to key **A**. If the print function is on and inputs are being printed, in both cases the printout will be of \dot{V} , the flow rate in ml/min. The number in the display at the end of routine **A** or **B** is \dot{V}_{corr} , the flow rate after correction for BSA. It is the size of this number that determines whether the standard or the maximum clearance will be calculated. This number will also be printed if the print function is on.

Equations:

$$\dot{V}(\text{ml/min}) = \frac{V(\text{ml})}{t(\text{min})}$$

$$\dot{V}_{\text{corr}} = \begin{cases} \frac{1.73}{\text{BSA}} \dot{V} & \text{if corrected for BSA} \\ \dot{V} & \text{if no correction for BSA} \end{cases}$$

Maximum clearance ($\dot{V}_{\text{corr}} > 2$):

$$C_m(\text{ml/min}) = \frac{U_{\text{urea}} \dot{V}_{\text{corr}}}{B_{\text{urea}}}$$

Standard clearance ($\dot{V}_{\text{corr}} \leq 2$):

$$C_s(\text{ml/min}) = \frac{U_{\text{urea}} \sqrt{\dot{V}_{\text{corr}}}}{B_{\text{urea}}}$$

where

U_{urea} = concentration of urea in urine

B_{urea} = concentration of urea in blood

$$\% \text{ mean normal } C_m = 1.33 C_m$$

$$\% \text{ mean normal } C_s = 1.85 C_s$$

Remarks:

1. Any units may be used for U_{urea} and B_{urea} as long as they are consistent.
2. Some users may prefer to ignore the distinction between standard and maximum clearance and use the maximum formula for all cases. This can be accomplished by using the program *Creatinine Clearance* (CL1-06A) and inputting U_{urea} and B_{urea} in place of U_{creat} and P_{creat} , respectively.
3. If the print function is turned off, neither inputs nor outputs will be printed.

Reference:

Clinical Chemistry, ed. Henry *et al.*, Harper and Row, 1974.

| STEP | INSTRUCTIONS | INPUT DATA/UNITS | KEYS | OUTPUT DATA/UNITS |
|------|--|------------------|----------|-------------------|
| 1 | Load side 1 and side 2. | | | |
| 2 | (optional) Initialize if reprint desired. | | A | 0.00 |
| 3 | (optional) Key in patient number. | Ptnt # | C | Ptnt # .05 |
| 4 | To suppress printing of data and results, turn the print function off. | | D | 0.00 |
| 5 | To turn the print function back on. | | D | 1.00 |

| STEP | INSTRUCTIONS | INPUT DATA/UNITS | KEYS | OUTPUT DATA/UNITS |
|------|---|-----------------------|-------|----------------------------------|
| 6 | If BSA is required and <i>Body Surface Area</i> has not been run, key in BSA. | BSA (m ²) | STO A | |
| 7 | If \dot{V} is to be corrected for BSA | | I B | BSA (m ²) |
| 8 | Perform either one of the steps below: | | | |
| | ● Key in urine volume and time | V (ml) | ENTER | |
| | | t (min) | A | \dot{V}_{corr} |
| | ● Key in urine flow rate | \dot{V} (ml/min) | B | \dot{V}_{corr} |
| 9 | Key in the concentration of urea in urine. | U _{urea} | C | U _{urea} |
| 10 | Key in the concentration of urea in blood and find the urea clearance. | B _{urea} | D | C _{urea} (ml/min) |
| 11 | Find the percent of mean normal. | | E | % m.n. |
| 12 | (optional) Reprint data and results. | | I E | Ptnt # .05 |
| | | | | \dot{V} |
| | | | | \dot{V}_{corr} |
| | | | | U _{urea} |
| | | | | B _{urea} |
| | | | | C _s or C _m |
| | | | | % m.n. |
| 13 | For a new case go to step 2. | | | |

Example 1:

A patient, number 10183, is to be tested for urea clearance. A volume of 204 ml of urine is collected over a period of 120 min. The concentration of urea in this urine is found to be 903 mg/100 ml. A blood sample is taken halfway through the urine collection and found to have a urea concentration of 26 mg/100 ml. Determine the urea clearance. Do not correct for body surface area.

Keystrokes:10183 **f** **C** →204 **ENTER** 120 **A** →903 **C** →26 **D** →**E** →**Outputs:**

10183.05 *** (Ptnt ID)

1.70 *** (\dot{V})1.70 *** (\dot{V}_{corr})903.00 *** (U_{urea})26.00 *** (B_{urea})45.28 *** (C_s , ml/min)

83.77 *** (% m.n.)

Example 2:

Patient number 10142 is a male, height 188 cm, weight 88.5 kg. A urine flow rate of 2.7 ml/min. is recorded. The concentration of urea is 798 mg/100 ml in urine and 21 mg/100 ml in blood. Determine the urea clearance corrected for body surface area using the Dubois formula for BSA.

Keystrokes:**f** **D** →188 **A** →88.5 **B** →**C** →**Outputs:**

0.00 (Print off)

188.00 (Ht, cm)

88.50 (Wt, kg)

2.15 (Dubois BSA)

Load side 1 and side 2 of *Urea Clearance* (CL1-05A).10142 **f** **C** →**f** **B** →2.7 **B** →798 **C** →21 **D** →**E** →

10142.05 *** (Ptnt ID)

2.15 (BSA)

2.70 *** (\dot{V})2.17 *** (\dot{V}_{corr})798.00 *** (U_{urea})21.00 *** (B_{urea})82.53 *** (C_m , ml/min)

109.76 *** (% m.n.)

CREATININE CLEARANCE



This program allows the calculation of creatinine clearance given the urine flow rate and the concentration of creatinine in urine and plasma. The urine flow rate may be corrected for the patient's body surface area if desired.

To indicate that a correction should be made for the body surface area, press **f** **B**. No action is necessary if the correction is not desired. If correction is to be made, the program will need to find the patient's body surface area (BSA, in m^2) in register R_A . The program *Body Surface Area* (CL1-04A) automatically leaves BSA stored in R_A . If *Body Surface Area* has not been run immediately before this program, you will need to key in the BSA and press **STO** **A**.

When inputting the urine flow rate, you may either key in the flow rate (\dot{V} , in ml/min.) directly to key **B**, or key in both the urine volume (V , in ml) and the time (t , in minutes) to key **A**. If the print function is on and inputs are being printed, in both cases the printout will be of \dot{V} , the flow rate in ml/min. The number in the display at the end of routine **A** or **B** is \dot{V}_{corr} , the flow rate after correction for BSA. (If no correction is desired, \dot{V}_{corr} will be the same as \dot{V} .) This number will also be printed if the print function is on.

Equations:

$$\dot{V}(\text{ml/min}) = \frac{V(\text{ml})}{t(\text{min})}$$

$$\dot{V}_{corr} = \begin{cases} \frac{1.73}{BSA} \dot{V} & \text{if corrected for BSA} \\ \dot{V} & \text{if not corrected for BSA} \end{cases}$$

$$C_{creat}(\text{ml/min}) = \frac{U_{creat} \dot{V}_{corr}}{P_{creat}}$$

where

C_{creat} = creatinine clearance

U_{creat} = concentration of creatinine in urine

P_{creat} = concentration of creatinine in plasma

Remarks:

- 1. Any units may be used for U_{creat} and P_{creat} as long as they are consistent.
- 2. If the print function is turned off, neither inputs nor outputs will be printed.

Reference:

Clinical Chemistry, ed. Henry *et al.*, Harper and Row, 1974.

| STEP | INSTRUCTIONS | INPUT DATA/UNITS | KEYS | OUTPUT DATA/UNITS |
|------|--|-----------------------|---------------------|-----------------------------|
| 1 | Load side 1 of program. | | | |
| 2 | (optional) Initialize if reprint | | | |
| | desired. | | f A | 0.00 |
| 3 | (optional) Key in patient | | | |
| | number. | Ptnt # | f C | Ptnt # .06 |
| 4 | To suppress printing of data | | | |
| | and results, turn print | | | |
| | function off. | | f D | 0.00 |
| 5 | To turn print function back | | | |
| | on later | | f D | 1.00 |
| 6 | If BSA is required and <i>Body</i> | | | |
| | <i>Surface Area</i> has not been | | | |
| | run, key in BSA. | BSA (m ²) | STO A | |
| 7 | If \dot{V} is to be corrected for BSA. | | f B | BSA (m ²) |
| 8 | Perform either one of the | | | |
| | steps below: | | | |
| | • Key in urine volume and | | | |
| | time | V (ml) | ENTER | |
| | | t (min) | A | \dot{V}_{corr} |
| | • Key in urine flow rate. | \dot{V} (ml/min) | B | \dot{V}_{corr} |
| 9 | Key in the concentration of | | | |
| | creatinine in urine. | U_{creat} | C | U_{creat} |
| 10 | Key in the concentration of | | | |
| | creatinine in plasma and find | | | |
| | the creatinine clearance. | P_{creat} | D | C_{creat} (ml/min) |

| STEP | INSTRUCTIONS | INPUT DATA/UNITS | KEYS | OUTPUT DATA/UNITS |
|------|--------------------------------------|---------------------|-------------------|-------------------------|
| 11 | (optional) Reprint data and results. | | f E | Ptnt # .06 |
| | | | | \dot{V} |
| | | | | \dot{V}_{corr} |
| | | | | U_{creat} |
| | | | | P_{creat} |
| | | | | C_{creat} |
| 12 | For a new case go to step 2. | | | |

Example 1:

A male, patient number 10095, is tested for creatinine clearance. A urine volume of 506 ml is collected over a 4-hour (240-min.) period. The concentration of creatinine is found to be 43.4 mg/dl in urine and 0.91 mg/dl in plasma. Find the creatinine clearance. Do not correct for body surface area.

Keystrokes:

10095 **f** **C** _____
506 **ENTER** 240 **A** _____
43.4 **C** _____
0.91 **D** _____

Outputs:

10095.06 *** (Ptnt ID)
2.11 *** (\dot{V})
2.11 *** (\dot{V}_{corr})
43.40 *** (U_{creat})
0.91 *** (P_{creat})
100.55 *** (C_{creat} , ml/min)

Example 2:

Patient number 10124 is a female with a body surface area of 1.56 m². Given a urine flow rate of 1.81 ml/min., a creatinine concentration of 46.5 mg/dl in urine and 1.03 mg/dl in plasma, find the creatinine clearance.

Keystrokes:

10124 **f** **C** _____
1.56 **STO** **A** _____
f **B** _____
1.81 **B** _____
46.5 **C** _____
1.03 **D** _____

Outputs:

10124.06 *** (Ptnt ID)
1.56 (BSA)
1.56
1.81 *** (\dot{V})
2.01 *** (\dot{V}_{corr})
46.50 *** (U_{creat})
1.03 *** (P_{creat})
90.62 *** (C_{creat} , ml/min)

Notes

AMNIOTIC FLUID ASSAY



AMNIOTIC FLUID ASSAY

CL1-07A

CLEAR
A₃₆₅

A₅₅₀

PTNT #
A₄₅₀ + Δ

P OFF?
Wk + b

REPRINT
Zone

This program performs calculations for the spectrophotometric estimation of bile pigments in amniotic fluid. Measurement of absorbance changes in the fluid has been shown to be useful in determining the management of Rh-sensitized pregnancies. The absorbance of the fluid is measured at two wavelengths (typically, 365 nm and 550 nm) to form a baseline, and then at a third wavelength between these two (typically, 450 nm) to allow calculation of the difference (Δ) between the actual and the interpolated absorbances at the intermediate wavelength. Then, given the weeks of gestation, the “b” factor and, optionally, the Liley zone number may be found.

The inputs to the program, then, are the absorbances of the amniotic fluid at three wavelengths (A₃₆₅, A₅₅₀, and A₄₅₀). From these may be found ΔA₄₅₀, the difference in absorbance at the intermediate wavelength. The final input is the week of gestation (Wk), from which may be found the “b” factor and zone. The last two outputs are the most meaningful for the obstetrician; for interpretation, see references 1 and 2 below.

Equations:

$$\Delta A_{450} = A_{450} - e^{[-.541 (\ln A_{365} - \ln A_{550}) + \ln A_{550}]}$$
$$b = \Delta A_{450} / a^{Wk}$$

where

a = 0.91509
Wk = week of gestation

Liley zones:

- Zone I: b < 0.7
- Zone II: 0.7 ≤ b ≤ 3
- Zone III: b > 3

Remarks:

1. Some users may prefer to take absorbance readings at wavelengths other than those indicated here. Burnett³, for instance, advocates readings at 350 nm, 550 nm, and 455 nm. It is quite easy to modify the program to handle such a case. The only change required is the alteration of

one constant occupying four steps of program memory, 024-027. At present in these locations the program holds the constant .541. For Burnett's values (350, 550, 455) this constant would have to be changed to .475. In general, if the three wavelengths used are x, y, and z, with $x < z < y$, the constant to be used is

$$\frac{y - z}{y - x}.$$

The absorbances at wavelengths x, y, and z should be input to keys **A**, **B**, and **C** respectively.

2. If the print function is turned off, neither inputs nor outputs will be printed.

References:

1. R.C. Brown and W.J. Beckfield, "Computer-assisted spectrophotometric analysis of amniotic fluid in erythroblastosis fetalis," *Amer. J. Clin. Path.*, **57**: 659-663, 1972.
2. A.W. Liley, "Liquor amnii analysis in the management of the pregnancy complicated by rhesus sensitization," *Amer. J. Obstet. Gynecol.*, **82**: 1359-1370, 1961.
3. R. Burnett, "Instrumental and procedural sources of error in determination of bile pigments in amniotic fluid," *Clin. Chem.*, **18**: 150-154, 1972.

| STEP | INSTRUCTIONS | INPUT DATA/UNITS | KEYS | OUTPUT DATA/UNITS |
|------|--|------------------|-------------------|-------------------|
| 1 | Load side 1 and side 2. | | | |
| 2 | (optional) Initialize if reprint desired. | | F A | 0.00 |
| 3 | (optional) Key in patient number. | Ptnt # | F C | Ptnt # .07 |
| 4 | To suppress printing of data and results, turn print function off. | | F D | 0.00 |
| 5 | To turn print function back on later. | | F D | 1.00 |
| 6 | Key in absorbance at 365 nm. | A ₃₆₅ | A | A ₃₆₅ |
| 7 | Key in absorbance at 550 nm. | A ₅₅₀ | B | A ₅₅₀ |

| STEP | INSTRUCTIONS | INPUT DATA/UNITS | KEYS | OUTPUT DATA/UNITS |
|------|--------------------------------|------------------|------------|-------------------|
| 8 | Key in absorbance at 450 nm | | | |
| | and find ΔA_{450} . | A_{450} | C | ΔA_{450} |
| 9 | Key in week of gestation and | | | |
| | find b factor. | Wk | D | b |
| 10 | (optional) Find Liley zone | | | |
| | number (1, 2, or 3). | | E | Zone |
| 11 | (optional) To obtain a reprint | | | |
| | of data and results. | | I E | Ptnt # .07 |
| | | | | A_{365} |
| | | | | A_{550} |
| | | | | A_{450} |
| | | | | ΔA_{450} |
| | | | | Week |
| | | | | b |
| | | | | Zone |

Example:

A sample of amniotic fluid from patient number 10070 is found to have absorbances of 0.43, 0.25, and 0.39 at wavelengths 365 nm, 550 nm, and 450 nm respectively. Find ΔA_{450} , the b factor, and the Liley zone number given that it is the 35th week of gestation.

Keystrokes:

10070 **I C** →
.43 **A** →
.25 **B** →
.39 **C** →

35 **D** →

E →

Outputs:

10070.07 *** (Ptnt ID)
0.43 *** (A_{365})
0.25 *** (A_{550})
0.39 *** (A_{450})
0.05 *** (ΔA_{450})
35. *** (Wk)
1.22 *** (b)
2. *** (Zone)

Notes

BLOOD ACID-BASE STATUS



This program computes total plasma CO_2 (TCO_2) and base excess (BE) from the partial pressure of CO_2 (PCO_2), pH, and hemoglobin concentration (Hgb). The PCO_2 and pH values used should be found at 37°C ; if they are found at a body temperature (BT) other than 37°C , the program will correct them to 37°C values if BT is also input. An additional, optional output of the program is the concentration of plasma bicarbonate ($[\text{HCO}_3^-]$).

To operate the program, if the body temperature is different from 37°C , then key in BT in $^\circ\text{C}$ and press key **A**. If $\text{BT} = 37^\circ\text{C}$, it need not be input; if it is, however, no harm will be done. Next key in PCO_2 in mm Hg and press **B**; the number displayed at the completion of this step is the value of PCO_2 corrected to 37°C . Then key in pH and press **C**; the result in the display at the end of this step is the pH value corrected to 37°C . Finally, press **D** to calculate TCO_2 in mmol/l. As an optional step, hemoglobin may now be input in units of g/100 ml. Pressing **E** will allow the calculation of base excess in mEq/l using an equation suggested by Siggaard-Andersen. The last value output is $[\text{HCO}_3^-]$, which may be found by pressing **R/S** after the calculation of base excess.

Equations:

$$\text{PCO}_2 (37^\circ\text{C}) = \text{PCO}_2 (\text{BT}) \cdot 10^{0.019 (37 - \text{BT})}$$

$$\text{pH} (37^\circ\text{C}) = \text{pH} (\text{BT}) - 0.0146 (37 - \text{BT})$$

$$\text{TCO}_2 = s \cdot \text{PCO}_2 [1 + 10^{\text{pH} - \text{pK}}]$$

where

s = solubility of CO_2 in plasma, mmol/l (taken to be 0.0307)

$\text{pK} = 6.11$

$$[\text{BE}]_b = (1 - 0.0143 \text{ Hgb}) ([\text{HCO}_3^-] - (9.5 + 1.63 \text{ Hgb}) (7.4 - \text{pH}) - 24)$$

where

$[\text{BE}]_b$ = base excess in mEq/l of blood

Hgb = hemoglobin concentration in g/100 ml

$$[\text{HCO}_3^-] = s \cdot \text{PCO}_2 \cdot 10^{\text{pH} - \text{pK}}$$

where






$[\text{HCO}_3^-]$ = concentration of plasma bicarbonate in mmol/l.

Remarks:

1. This program can also be used to correct PCO_2 and pH values from 37°C to body temperature. To do this, let $x = (74 - \text{BT})^\circ\text{C}$. Key in x to key **A**. Then input PCO_2 and pH to keys **B** and **C**, respectively. The number displayed after each of these steps is the value of the parameter corrected to body temperature. For example, if it is desired to correct a 37°C PCO_2 value of 45 mm Hg to a body temperature value with $\text{BT} = 40^\circ\text{C}$, let $x = 34$. Key in 34, press **A**, key in 45, and press **B**. The corrected PCO_2 is found to be 51.31 mm Hg.
2. The equation to correct pH to 37°C values is a simplification of a formula from Severinghaus. It ignores the pH and BE dependent terms. This introduces a very small error except at extreme conditions of acid-base status and large temperature shifts. For example, at a pH of 7.2 or 7.6, the error is 0.0013 units per $^\circ\text{C}$.
3. If the print function is turned off, neither inputs nor outputs will be printed.

References:

- John W. Severinghaus, "Blood gas calculator," *J. Appl. Physiol.*, **21**: 1108 – 1116, 1966.
- Siggaard-Andersen, "Titrable acid or base of body fluids," *Annals New York Academy of Science*, **133**: 41-48, 1966.
- L.J. Thomas, Jr., "Algorithms for selected blood acid-base and blood gas calculation," *J. Appl. Physiol.*, **33**: 154-158, 1972.

| STEP | INSTRUCTIONS | INPUT DATA/UNITS | KEYS | OUTPUT DATA/UNITS |
|------|---|--------------------------|--|--|
| 1 | Load side 1 and side 2 of program. | | | |
| 2 | (optional) Initialize if reprint desired. | |  A | 0.00 |
| 3 | (optional) Key in patient number. | Ptnt # |  C | Ptnt # .08 |
| 4 | To suppress printing of data and results, turn print function off. | |  D | 0.00 |
| 5 | To turn print function back on later. | |  D | 1.00 |
| 6 | If PCO ₂ and pH are to be corrected to 37°C, key in body temperature in °C. | BT (°C) | A | 37 – BT |
| 7 | Key in partial pressure of CO ₂ in mm Hg. | PCO ₂ (mm Hg) | B | PCO ₂ (37°) |
| 8 | Key in pH. | pH | C | pH (37°) |
| 9 | Find total plasma CO ₂ in mmol/l. | | D | TCO ₂ (mmol/l) |
| 10 | (optional) Key in hemoglobin concentration and compute base excess and [HCO ₃ ⁻] | Hgb(g/100ml) | E | BE (mEq/l) |
| | | | R/S | [HCO ₃ ⁻](mmol/l) |
| 11 | To obtain a reprint. | |  E | Ptnt # .08 |
| | | | | BT |
| | | | | PCO ₂ |
| | | | | pH |
| | | | | TCO ₂ |
| | | | | Hgb |
| | | | | BE |
| | | | | [HCO ₃ ⁻] |

Example :

Patient number 10183 has a body temperature of 40°C. His PCO₂ at 40°C is found to be 51 mm Hg, his pH at the same temperature 7.31. His hemoglobin concentration is 16 g/100 ml. Find TCO₂, BE, and [HCO₃⁻].

Keystrokes:

f **A** _____→
10183 **f** **C** _____→
40 **A** _____→

51 **B** _____→

7.31 **C** _____→

D _____→
16 **E** _____→

R/S _____→

Outputs:.

0.00
10183.08 *** (Ptnt ID)
40.00 *** (BT)
-3.00 (37 - BT)
51.00 *** (PCO₂(40))
44.73 (PCO₂(37))
7.31 *** (pH(40))
7.35 (pH(37))
25.44 *** (TCO₂)
16.00 *** (Hgb)
-1.21 *** (BE)
24.07 *** ([HCO₃⁻])

OXYGEN SATURATION AND CONTENT



This program estimates oxygen saturation of blood from various body parameters and computes oxygen content. If the actual oxygen saturation is known, oxygen content may be computed directly.

Estimated saturation

Typically, the input parameters to the program are PCO_2 , pH, and PO_2 measured at 37°C , and the body temperature in $^\circ\text{C}$. If the parameters PCO_2 and pH are known only at body temperature, they may be corrected to 37°C through use of the program *Blood Acid-Base Status*, CL1-08A. If CL1-08A is run before this program, the values of BT, PCO_2 , and pH may be recalled by this program for input to the appropriate keys. For example, pressing **f** **B** will recall the value of BT. Pressing **A** will then input the recalled value to this program and recall the value of PCO_2 . Pressing **B** will input the recalled PCO_2 value and recall the value of pH. If CL1-08A has not been run previously, the recalled values will be meaningless numbers or zero.

After the input of PO_2 to **D**, an intermediate value of virtual PO_2 (VPO_2) will be calculated prior to the calculation of estimated saturation. The value found for VPO_2 will not be output but may be displayed after the calculation of saturation by pressing **RCL C**. VPO_2 is not a real physiologic PO_2 . Its only use is in estimating O_2 saturation.

Suppose as an alternate case that BT, PCO_2 , and pH are not known, but virtual PO_2 , or alveolar PO_2 ($\text{P}_\text{A}\text{O}_2$) is known. In this case, only the known VPO_2 or $\text{P}_\text{A}\text{O}_2$ need be input in order to compute estimated saturation. Input VPO_2 or $\text{P}_\text{A}\text{O}_2$ to key **D** as *negative* numbers, i.e., key in the value followed by **CHS**, then press **D**. The output, as before, will be estimated oxygen saturation.

After computing saturation, the hemoglobin concentration in g/100 ml should be keyed into **E**. Output from this sequence will be the oxygen content as a volume percent.

Known saturation

If the actual O_2 saturation is known, the oxygen content may be computed directly. Simply key in the O_2 saturation, press **ENTER**, key in hemoglobin concentration and press **E**. Oxygen content will be output.

Equations:

$$\text{VPO}_2 = \text{PO}_2 \cdot 10^{[0.024(37-\text{BT}) + 0.48(\text{pH}-7.4) + 0.06 \log(40/\text{PCO}_2)]}$$

$$\text{O}_2 \text{ Sat} = \frac{(\text{VPO}_2)^4 - 15(\text{VPO}_2)^3 + 2045(\text{VPO}_2)^2 + 2000(\text{VPO}_2)}{(\text{VPO}_2)^4 - 15(\text{VPO}_2)^3 + 2400(\text{VPO}_2)^2 - 31,100(\text{VPO}_2) + 2,400,000}$$






$$\text{O}_2 \text{ content} = 1.34 \cdot \frac{\text{Sat} (\%)}{100} \cdot \text{Hgb} + 0.0031 \text{ VPO}_2$$

Remarks:

1. In the computation of VPO_2 , it is important to input the values for pH and BT exactly, as these have a great influence on the value of VPO_2 . PCO_2 has relatively little influence.
2. The equation for VPO_2 is a hybrid of the equation used by Thomas and that used by Kelman. There is some disagreement regarding the best value of the pH multiplier, 0.48 being used by most workers, but see, for example, Kelman.
3. The calculation of saturation from PO_2 will give inaccurate results for fetal hemoglobin, present in babies less than six months old, and for some abnormal adult hemoglobins and certain other blood conditions. The results of the estimation and any subsequent calculations based on it, should be viewed with caution unless the dissociation curve has been previously established to be normal. If both PO_2 and O_2 saturation are measured, the program may be used as a convenient means to check for the normality of the dissociation curve.
4. If the print function is turned off, neither inputs nor outputs will be printed.
5. After a keystroke sequence in which **D** is pressed to find saturation, **E** should also be pressed to complete the sequence even if Hgb is not input and the calculated oxygen content is meaningless.

References:

- L.J. Thomas, Jr., "Algorithms for selected blood acid-base and blood gas calculation," *J. Appl. Physiol.*, **33**: 154-158, 1972.
- G. Richard Kelman, "Digital computer subroutine for the conversion of oxygen tension into saturation," *J. Appl. Physiol.*, **21**: 1375-1376, 1966.

| STEP | INSTRUCTIONS | INPUT DATA/UNITS | KEYS | OUTPUT DATA/UNITS |
|------|---|--------------------------|--|------------------------------|
| 1 | Load side 1 and side 2. | | | |
| 2 | (optional) Initialize if reprint desired. | |  A | 0.00 |
| 3 | (optional) Key in patient number. | Ptnt # |  C | Ptnt # .09 |
| 4 | To suppress printing of data and results, turn print function off. | |  D | 0.00 |
| 5 | To turn print function back on later. | |  D | 1.00 |
| 6 | If oxygen saturation is to be estimated, go to step 7; if it is known already, go to step 14. | | | |
| | Estimated saturation | | | |
| 7 | If BT was stored from <i>Blood Acid-Base Status</i> (CL1-08A), it may be recalled. | |  B | BT (°C) |
| 8 | Input body temperature in °C. | BT (°C) | A | PCO ₂ (if stored) |
| 9 | Input PCO ₂ in mm Hg. | PCO ₂ (mm Hg) | B | pH(if stored) |
| 10 | Input pH. | pH | C | pH |
| 11 | Input PO ₂ in mm Hg (CHS for VPO ₂ or P _A O ₂) and find oxygen saturation. | PO ₂ (mm Hg) | D | Sat (%) |
| 12 | Key in hemoglobin and find oxygen content as a volume percent. | Hgb (g/100ml) | E | O ₂ content |

| STEP | INSTRUCTIONS | INPUT DATA/UNITS | KEYS | OUTPUT DATA/UNITS |
|------|------------------------------|------------------|-------------------|------------------------|
| 13 | To obtain a reprint. | | f E | Ptnt # .09 |
| | | | | BT |
| | | | | PCO ₂ |
| | | | | pH |
| | | | | PO ₂ |
| | | | | Sat (%) |
| | | | | Hgb |
| | | | | O ₂ content |
| | Known saturation | | | |
| 14 | Key in saturation and | | | |
| | hemoglobin concentration and | | | |
| | find oxygen content as a | | | |
| | volume percent. | Sat (%) | ENTER | |
| | | Hgb (g/100ml) | E | O ₂ content |

Example 1:

Patient number 10183 has a body temperature of 40°C. The following parameters are measured at 37°C: PCO₂ = 45 mm Hg, pH = 7.35, and PO₂ = 75 mm Hg. Find the estimated O₂ saturation. Given a hemoglobin concentration of 16 g/100 ml, find oxygen content.

Keystrokes:

10183 **f** **C** →
40 **A** →
45 **B** →
7.35 **C** →
75 **D** →
16 **E** →

Outputs:

10183.09 *** (Ptnt ID)
40.00 *** (BT)
45.00 *** (PCO₂)
7.35 *** (pH)
75.00 *** (PO₂)
90.92 *** (Sat %)
16.00 *** (Hgb)
19.68 *** (O₂ cont.)

Example 2:

Alveolar PO₂ (P_A O₂) is known to be 103 mm Hg in patient number 10184. Find the estimated O₂ saturation. Given a hemoglobin concentration of 14.5 g/100 ml, find the oxygen content.

Keystrokes:

10184 **f** **C** →
103 **CHS** **D** →
14.5 **E** →

Outputs:

10184.09 *** (Ptnt ID.)
-103.00 *** (P_AO₂)
97.72 *** (Sat %)
14.50 *** (Hgb)
19.31 *** (O₂ cont.)

Example 3:

Oxygen saturation is measured at 92%. Hemoglobin concentration is 16 g/100 ml. What is the oxygen content?

92 **ENTER** 16 **E** →

92.00 *** (Sat %)
16.00 *** (Hgb)
20.04 *** (O₂ cont.)

Notes

RED CELL INDICES



This program computes red cell indices based on three measured values: red cell count, hematocrit, and hemoglobin. The indices computed are mean corpuscular volume (MCV), mean corpuscular hemoglobin (MCH), and mean corpuscular hemoglobin concentration (MCHC).

The red cell count in millions per mm³ should be input to key **A** and hematocrit as a percent to key **B**. Then hemoglobin in g/dl (g/100 ml) is keyed in, and **C** is pressed to allow calculation of MCV in cubic microns (μ³). Pressing **D** will cause the output of MCH in picograms, pg (or micromicrograms, μμg). Finally, key **E** is pressed to compute MCHC in g/dl (g/100 ml).

Equations:

$$MCV (\mu^3) = \frac{Hct (\%) \times 10}{Count (10^6/mm^3)}$$

$$MCH (pg) = \frac{Hgb (g/dl) \times 10}{Count (10^6/mm^3)}$$
















$$MCHC (g/dl) = \frac{Hgb (g/dl) \times 100}{Hct (\%)}$$

Remarks:

If the print function is turned off, neither inputs nor outputs will be printed.

Reference:

Davidson and Henry, *Todd-Sanford Clinical Diagnosis by Laboratory Methods*, W.B. Saunders Co., 1969.

| STEP | INSTRUCTIONS | INPUT DATA/UNITS | KEYS | OUTPUT DATA/UNITS |
|------|---|---|---|-----------------------|
| 1 | Load side 1. | | | |
| 2 | (optional) Initialize if reprint desired. | |   | 0.00 |
| 3 | (optional) Key in patient number. | Ptnt # |   | Ptnt # .10 |
| 4 | To suppress printing of data and results, turn print function off. | |   | 0.00 |
| 5 | To turn print function back on later. | |   | 1.00 |
| 6 | Key in red cell count in millions per mm ³ . | Count (10 ⁶ /mm ³) |  | Count |
| 7 | Key in hematocrit. | Hct (%) |  | Hct (%) |
| 8 | Key in hemoglobin in g/100ml and find mean corpuscular volume in μ ³ . | Hgb (g/dl) |  | MCV (μ ³) |
| 9 | Compute mean corpuscular hemoglobin in pg (μμg). | |  | MCH (pg) |
| 10 | Compute mean corpuscular hemoglobin concentration in g/dl (g/100ml). | |  | MCHC (g/dl) |
| 11 | To obtain a reprint of data and results. | |   | Ptnt # .10 |
| | | | | Count |
| | | | | Hct (%) |
| | | | | Hgb |
| | | | | MCV |
| | | | | MCH |
| | | | | MCHC |

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Example:

A sample of venous blood from patient 10183 reveals a red cell count of $2.25 \times 10^6/\text{mm}^3$, a hematocrit of 21%, and hemoglobin of 7.2 g/dl (g/100 ml). Find the indices MCV, MCH, and MCHC.

Keystrokes:

10183 **f** **C** →
2.25 **A** →
21 **B** →
7.2 **C** →

D →
E →

Outputs:

10183.10 *** (Ptnt ID)
2.25 *** (Count)
21.00 *** (Hct %)
7.20 *** (Hgb)
93.33 *** (MCV)
32.00 *** (MCH)
34.29 *** (MCHC)

Notes

TOTAL BLOOD VOLUME



This program computes total blood volume by the radioisotope dilution technique. The inputs to the program are the background counts per minute (Bck), the volume of radioactive solution injected (V Inj), the dilution of the standard solution (Std Dil), the counts per minute of the standard (Std CPM), and the counts per minute of the sample of whole blood (WB CPM). From these values the program will compute total blood volume (TBV).

Equations:

$$TBV = Dil \times V\ Inj \times \frac{Std\ CPM - Bck}{WB\ CPM - Bck}$$

Remarks:

- 1. Total blood volume will be computed in the same units as volume injected. Typically the units used will be milliliters (ml).
- 2. Equal volumes of whole blood, diluted standard solution, and distilled water should be used for the measurement of whole blood counts, standard counts, and background counts. These three counts need not be counts *per minute*; they may be counts recorded over any length of time, so long as the same time interval is used for all three counts.
- 3. This same program may be used to find total plasma volume provided that a sample of plasma rather than whole blood is counted for the final input. Total blood volume may be determined from total plasma volume from the equation

$$Total\ blood\ volume = \frac{Total\ plasma\ volume}{(1 - Hct \times 0.9)}$$

- 4. If the patient has had prior radioactivity administered, a patient back-ground correction may be necessary. To do this, a count must be made of a blood sample before the current dose is administered. These pre-dose counts should be subtracted from the post-dose whole blood counts to give the corrected counts to be input at the final step.
- 5. If the print function is turned off, neither inputs nor outputs will be printed.

Reference:

Beierwaltes, Keyes, and Carey, *Manual of Nuclear Medicine Procedure*, Chemical Rubber Co., 1971.

| STEP | INSTRUCTIONS | INPUT DATA/UNITS | KEYS | OUTPUT DATA/UNITS |
|------|--|------------------|------------|-------------------|
| 1 | Load side 1. | | | |
| 2 | (optional) Initialize for reprint. | | F A | 0.00 |
| 3 | (optional) Key in patient number. | Ptnt # | F C | Ptnt # .11 |
| 4 | To suppress printing of data and results, turn print function off. | | F D | 0.00 |
| 5 | To turn print function back on later. | | F D | 1.00 |
| 6 | Key in background counts. | Bck | A | Bck |
| 7 | Key in volume of fluid injected. | Vol. inj. | B | Vol. inj. |
| 8 | Key in dilution of standard. | Std. dil. | C | Std. dil. |
| 9 | Key in standard counts. | Std. CPM | D | Std. CPM |
| 10 | Key in whole blood counts and find total blood volume. | Blood CPM | E | TBV |
| 11 | To obtain a reprint. | | F E | Ptnt # .11 |
| | | | | Bck |
| | | | | Vol. inj. |
| | | | | Std. dil. |
| | | | | Std. CPM |
| | | | | Blood CPM |
| | | | | TBV |

Example:

5 ml of radioiodinated serum albumin (RISA) are injected into patient 10183. The stock RISA is diluted by a factor of 250 and a 1 ml aliquot of this standard is found to have an activity of 2518 counts over a five-minute period. A 1 ml sample of the patient's whole blood, collected 10 minutes after injection, is found to have an activity of 837 counts over a five-minute period. A five-minute count of 1 ml distilled water yields 152 counts. What is the patient's total blood volume?

Keystrokes:

10183 **f** **C** →
152 **A** →
5 **B** →
250 **C** →
2518 **D** →
837 **E** →

Outputs:

10183.11 *** (Ptnt ID)
152.00 *** (Bck)
5.00 *** (V Inj)
250.00 *** (Dil)
2518.00 *** (Std CPM)
837.00 *** (WB CPM)
4317.52 *** (TBV, ml)

Notes

SCHILLING TEST



This program performs the calculations involved with the Schilling test for the determination of vitamin B₁₂ absorption. The inputs to the program are the background counts per minute, the dilution and counts per minute of the standard, the volume of urine excreted, and the counts per minute of the urine. The output is the % of dose excreted.

The program is set up to handle urine volume (U Vol) in liters (l). It is assumed that if the urine volume collected was less than 1 l, the volume was brought up to 1 l by the addition of water. If the volume was a liter or more, no dilution should be made.

Equations:

$$\% \text{ excretion} = \frac{V}{\text{Dil}} \left[\frac{\text{Urine CPM} - \text{Background CPM}}{\text{Standard CPM} - \text{Background CPM}} \right] \times 100$$

$$\text{where } V = \begin{cases} 1 & \text{if } U \text{ Vol} \leq 1 \text{ l} \\ U \text{ Vol} & \text{if } U \text{ Vol} > 1 \text{ l} \end{cases}$$
















Dil = Dilution of the standard

Remarks:

- 1. The background, standard, and urine counts should be of equal volumes counted over equal time intervals (which need not be one minute).
- 2. The patient should not have had recent prior radioactivity.
- 3. If the print function is turned off, neither data nor results will be printed.

Reference:

Beierwaltes, Keyes, and Carey, *Manual of Nuclear Medicine Procedures*, Chemical Rubber Co., 1971.







| STEP | INSTRUCTIONS | INPUT DATA/UNITS | KEYS | OUTPUT DATA/UNITS |
|------|--|------------------|---|-------------------|
| 1 | Load side 1. | | | |
| 2 | (optional) Initialize if reprint is desired. | |   | 0.00 |
| 3 | (optional) Key in patient number. | Ptnt # |   | Ptnt # .12 |
| 4 | To suppress printing of data and results, turn print function off. | |   | 0.00 |
| 5 | To turn print function back on later. | |   | 1.00 |
| 6 | Key in background counts. | Bck |  | Bck |
| 7 | Key in dilution of the standard. | Std Dil |  | Std Dil |
| 8 | Key in standard counts. | Std CPM |  | Std CPM |
| 9 | Key in volume of urine collected. | U Vol (l) |  | U Vol |
| 10 | Key in the urine counts and calculate percentage of dose excreted. | U CPM |  | % |
| 11 | To obtain a reprint of data and results. | |   | Ptnt # .12 |
| | | | | Bck |
| | | | | Std Dil |
| | | | | Std CPM |
| | | | | U Vol |
| | | | | U CPM |
| | | | | % |

Example:

A capsule of radioactive B_{12} is administered orally to patient 10183. Over the following 24 hours, a volume of 2.54 l of urine is collected. A 20 ml aliquot of the urine is counted for 10 minutes to give 1923 counts. A 1 ml sample of the standard is diluted to 20 ml and counted for 10 minutes, giving 1757 counts. 20 ml of tap water is used for a background count; over a ten-minute interval, 127 counts are recorded. Find the percent of dose excreted.

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Keystrokes:

10183   →
127  →
20  →
1757  →
2.54  →
1923  →

Outputs:

10183.12 *** (Ptnt ID)
127.00 *** (Bck)
20.00 *** (Std Dil)
1757.00 *** (Std CPM)
2.54 *** (U Vol)
1923.00 *** (U CPM)
13.99 *** (% excreted)

Notes

THYROID UPTAKE



This program computes thyroid uptake as a percentage of an administered dose of radioiodine. The inputs to the program are the counts per minute for the standard, the standard background, the patient counts (after ingestion of the dose), and the patient background. After these variables have been input, pressing **E** will allow computation of the percent uptake.

After calculation of the uptake, two corrections may be made to the computed value. The first correction involves recent prior radioactivity in the patient. The second correction involves a significant difference in activity between the standard and the dose. These are discussed in more detail below.

If the patient has had recent prior radioactivity, the computed uptake must be corrected to account for this. In such a case the patient counts and the background counts *before* ingestion of the present dose must be known. In addition, it will be necessary to correct these predose counts for radioactive decay over the elapsed time between the measurements of the predose counts and of the counts after ingestion of the dose. The program *Radioactive Decay Corrections* (CL1-14A) may be used to account for this decay. *Radioactive Decay Corrections* will compute and store a decay factor *D* that will be used by this program, *Thyroid Uptake*, to adjust the predose counts to the present time.

To correct for prior radioactivity, then, you should first load side 1 and side 2 of *Radioactive Decay Corrections* (CL1-14A). Select the radioisotope of the *prior* radioactivity. Key in 1, press **A**, then key in the time interval over which the decay has occurred, in the format DD.HH (days.hours), remembering always to allow 2 places for hours. (For example, a period of 1 day 6 hours should be keyed in as 1.06.) After keying in the elapsed time, press **B**, then press **C**. The decay factor *D* will be displayed and automatically stored. Now load side 1 and side 2 of *Thyroid Uptake* and follow the basic procedure to find the uncorrected percentage uptake. After computing % uptake from key **E**, key in the predose patient counts, press **ENTER**, key in the predose background counts and press **f** **A**. The corrected percentage uptake will be computed.

The second possible correction to be made is to account for a significant difference in the activities of the standard and the dose. These activities should be measured before the dose is administered. The counts at this point are referred to as precounts. If the standard and dose precounts agree within $\pm 3\%$, no correction is necessary. If the precounts differ by more than 3%, however, then the computed thyroid uptake should be corrected. To make the correction, after pressing **E** to find the uptake, key in the standard precount,

press **ENTER**, key in the dose precount, and press **f B**. The program will compute the corrected thyroid uptake.

The two corrections to computed uptake operate independently of each other. Either, both, or neither correction may be made. If both are to be made, they may be made in either order. If a reprint is called for after a correction is made, the reprint will show the corrected value of uptake but will not show the inputs that went into the correction (i.e., the patient and background pre-dose counts or the standard and dose precounts).

Equations:

$$\% \text{ uptake} = K \times \frac{\text{NPC}}{\text{Std CPM} - \text{Std Bck}} \times 100$$

where

$$\begin{aligned} \text{NPC} &= \text{Net Ptnt Cts} \\ &= \text{Ptnt CPM} - \text{Ptnt Bck} \end{aligned}$$

and K is a correction factor.

$$K = \begin{cases} 1 & \text{if no correction} \\ \frac{\text{NPC} - D \times (\text{Ptnt Predose Ct} - \text{Bck Predose Ct})}{\text{NPC}} & \text{if prior radioactivity} \\ \frac{\text{Std. Precount}}{\text{Dose Precount}} & \text{if different activities} \end{cases}$$

where

D is the radioactive decay factor.

Remarks:

1. The counts need not be input as counts *per minute*; however, all counts should be measured over the same time interval.
2. If the print function is turned off, neither inputs nor outputs will be printed.

Reference:

Beierwaltes, Keyes, and Carey, *Manual of Nuclear Medicine Procedures*, Chemical Rubber Co., 1971.

| STEP | INSTRUCTIONS | INPUT DATA/UNITS | KEYS | OUTPUT DATA/UNITS |
|------|--|------------------|------------|-------------------|
| 1 | If correction is to be made for prior patient radio-activity, go to step 2. Otherwise go to step 6. | | | |
| 2 | Load side 1 and side 2 of <i>Radioactive Decay Corrections</i> (CL1-14A) and select the radioisotope of the prior radioactivity. | | | |
| 3 | Key in a 1 for the initial activity. | 1 | A | 1.00 |
| 4 | Key in time elapsed in format Days.Hours (e.g., 1 day 6 hours is keyed in as 1.06). | t(dd.hh) | B | t(dd.hh) |
| 5 | Compute the decay factor (will be stored automatically). | | C | D |
| | Basic Procedure | | | |
| 6 | Load side 1 and side 2 of <i>Thyroid Uptake</i> (CL1-13A). | | | |
| 7 | (optional) Key in patient number. | Ptnt # | F C | Ptnt # .13 |
| 8 | To suppress printing of data and results, turn print function off. | | F D | 0.00 |
| 9 | To turn print function back on later. | | F D | 1.00 |
| 10 | Key in counts for the standard. | Std. CPM | A | Std. CPM |
| 11 | Key in background counts for the standard. | Std Bck. | B | Net Std. Cts. |
| 12 | Key in counts for the patient. | Ptnt. CPM | C | Ptnt. CPM |

| STEP | INSTRUCTIONS | INPUT DATA/UNITS | KEYS | OUTPUT DATA/UNITS |
|------|--|------------------|--------------|-------------------|
| 13 | Key in background counts for the patient. | Ptnt. Bck. | D | Net Ptnt. Cts. |
| 14 | Compute thyroid uptake as a percent. | | E | % Uptake |
| | Corrections | | | |
| 15 | For prior radioactivity, go to step 16, for differences in standard and dose, go to step 19. For no correction, go to step 20. | | | |
| | Prior Radioactivity | | | |
| 16 | For prior radioactivity, CL1-14A should have been run at step 2. | | | |
| 17 | Now key in patient predose counts and predose background and compute the corrected percent uptake. | Predose Cts. | ENTER | |
| | | Predose Bck. | I A | % Uptake |
| 18 | For differences in dose and standard, go to step 19. Otherwise go to step 20. | | | |
| | Differences in dose and standard | | | |
| 19 | Key in standard and dose precounts and find the corrected percent uptake. | Std. Prec. | ENTER | |
| | | Dose Prec. | I B | % Uptake |

| STEP | INSTRUCTIONS | INPUT DATA/UNITS | KEYS | OUTPUT DATA/UNITS |
|------|-----------------------------|------------------|-------------------|-------------------|
| | Reprint | | | |
| 20 | To obtain a reprint of data | | | |
| | and results. | | f E | Ptnt # .13 |
| | | | | Std. CPM |
| | | | | Std. Bck. |
| | | | | Ptnt. CPM |
| | | | | Ptnt. Bck. |
| | | | | % Uptake |

Example 1:

Before a dose of radioiodine (¹³¹I) is administered to patient 10183, a count is made of the patient’s current level of radioactivity from a prior ingestion of ¹³¹I. The patient’s predose activity is found to be 75 counts per minute (CPM) and the background predose activity 25 CPM. Twenty-four hours after ingestion of the dose, the patient’s activity is measured as 350 CPM with a background of 100 CPM. The activity of a standard of ¹³¹I is measured at 1500 CPM with a background of 200 CPM. Find the percentage uptake corrected for prior radioactivity.

Keystrokes:

Outputs:

Load side 1 and side 2 of *Radioactive Decay Corrections* (CL1-14A).

Select ¹³¹I as prior radioisotope.

| | | |
|------------------------|--------|-------------------------------|
| f E → | 193.20 | (¹³¹ I half-life) |
| 1 A → | 1.00 | |
| 0.24 B → | 0.24 | (24 hours) |
| C → | 0.92 | (Decay factor) |

Load side 1 and side 2 of *Thyroid Uptake* (CL1-13A).

| | | |
|--|--------------|--------------------|
| 10183 f C → | 10183.13 *** | (Ptnt ID) |
| 1500 A → | 1500.00 *** | (Std CPM) |
| 200 B → | 200.00 *** | (Std Bck) |
| | 1300.00 | (Net Std CPM) |
| 350 C → | 350.00 *** | (Ptnt CPM) |
| 100 D → | 100.00 *** | (Ptnt Bck) |
| | 250.00 | (Net Ptnt CPM) |
| E → | 19.23 *** | (% uptake) |
| 75 ENTER 25 f A → | 75.00 *** | (Ptnt Predose) |
| | 25.00 *** | (Bck Predose) |
| | 15.70 *** | (Corrected uptake) |

Example 2:

A standard and a dose are measured (before ingestion of the dose) at activities of 14,500 and 12,500 counts. Since the activities differ by more than 3%, a correction will have to be made to the computed percentage uptake. After ingestion of the dose, the standard activity is found to be 11,500 counts with a background count of 1000. The patient's activity is found as 2650 counts with a background of 500 counts. Find the corrected uptake.

Keystrokes:

11500 **A** →
1000 **B** →

2650 **C** →
500 **D** →

E →
14500 **ENTER** 12500 **f** **B** →

Outputs:

11500.00 *** (Std Cts)
1000.00 *** (Std Bck)
10500.00 (Net Std Cts)
2650.00 *** (Ptnt CPM)
500.00 *** (Ptnt Bck)
2150.00 (Net Ptnt Cts)
20.48 *** (% Uptake)
14500.00 *** (Std Prects)
12500.00 *** (Dose Prects)
23.75 *** (Corrected uptake)

RADIOACTIVE DECAY CORRECTIONS



This program is designed to allow calculation of the decay in radioactivity of an isotope over a specified time interval. The half-lives of 15 different radioisotopes are stored by the program and may be used in calculating the decay. Generally, to use the program you will select an isotope, key in the activity A_0 at the initial time, then key in the elapsed time t and calculate the present activity A . There are thus three variables needed to define the problem entirely: A_0 , t , and A .

An additional feature of the program is its ability to calculate *any* one of these variables given the other two. Thus you are not restricted to finding the present activity given the initial activity and time; you may also solve for initial activity given time and present activity, or for time given initial activity and present activity.

The radioisotope to be selected must be specified in one of two ways. Six isotopes are available directly by pressing user-definable keys **E** and **f A** through **f E**. Nine additional isotopes are available by keying in a digit, 1 through 9, and pressing **D**. For instance, to specify use of the radioisotope ^{57}Co , simply press **f B**. To specify the isotope ^{14}C , key in the number 2 and press **D**. A table of the correspondence between the isotopes and the numbers 1-9 may be found in the User Instructions. A list of available isotopes and their assumed half-lives is shown below.

You may use any units for the initial and present radioactivity, so long as they are consistent. The elapsed time must be input in the units Days. Hours (DD.HH), where two full decimal places must be allotted to the hours. For instance, an elapsed time of 5 days 18 hours would be keyed in and displayed as 5.18; a time of 1 day 6 hours as 1.06; and a time of 12 hours as 0.12.

Equations:

$$A = A_0 \left(\frac{1}{2} \right)^{t/\tau_{1/2}}$$

$$t = \frac{\tau_{1/2} \ln (A/A_0)}{\ln (1/2)}$$

where:

A_0 = initial radioactivity

A = present radioactivity

t = time elapsed, in hours

$\tau_{1/2}$ = half-life of radioisotope, in hours

| Isotope | $\tau_{1/2}$ (hrs) |
|---------------------------|---------------------|
| ^{51}Cr | 667.2 |
| ^{57}Co | 6480 |
| $^{99\text{m}}\text{Tc}$ | 6 |
| ^{125}I | 1440 |
| ^{131}I | 193.2 |
| ^{137}Cs | 262980 |
| ^3H | 107470 |
| ^{14}C | 5.058×10^7 |
| ^{18}F | 1.87 |
| ^{32}P | 343.2 |
| ^{75}Se | 2880 |
| ^{85}Sr | 1536 |
| $^{113\text{m}}\text{In}$ | 1.73 |
| ^{133}Xe | 126.5 |
| ^{197}Hg | 65 |

Remarks:

1. It is also possible to use this program for isotopes other than those provided by the program. In such a case, instead of selecting a radioisotope by the usual means, simply key in half-life in hours of the new isotope and press **STO** **B**. Then execute the rest of the program in the same fashion as usual.
2. Hours are not always rounded nicely to days for output. For example, a time of 6 days 23.8 hours would be computed in days. hours format as 6.238. In display mode FIX DSP 2, this would appear as 6.24, even though 7.00 might be the preferred rounded format.
3. Neither inputs nor outputs will be printed by the program.

| STEP | INSTRUCTIONS | INPUT DATA/UNITS | KEYS | OUTPUT DATA/UNITS |
|------|---------------------------------------|---------------------|-------------------|----------------------|
| 1 | Load side 1 and side 2. | | | |
| 2 | Select one of the fifteen radio- | | | |
| | isotopes and display half-life | | | |
| | in hours: | | | |
| | ● Chromium—51 (⁵¹ Cr) | | I A | 667.20 |
| | ● Cobalt—57 (⁵⁷ Co) | | I B | 6480.00 |
| | ● Technetium—99m (^{99m} Tc) | | I C | 6.00 |
| | ● Iodine—125 (¹²⁵ I) | | I D | 1440.00 |
| | ● Iodine—131 (¹³¹ I) | | I E | 193.20 |
| | ● Cesium—137 (¹³⁷ Cs) | | E | 262980.00 |
| | ● Hydrogen—3 (³ H) | 1 | D | 107470.00 |
| | ● Carbon—14 (¹⁴ C) | 2 | D | 50580000.00 |
| | ● Flourine—18 (¹⁸ F) | 3 | D | 1.87 |
| | ● Phosphorus—32 (³² P) | 4 | D | 343.20 |
| | ● Selenium—75 (⁷⁵ Se) | 5 | D | 2880.00 |
| | ● Strontium—85 (⁸⁵ Sr) | 6 | D | 1536.00 |
| | ● Indium—113m (^{113m} In) | 7 | D | 1.73 |
| | ● Xenon—133 (¹³³ Xe) | 8 | D | 126.50 |
| | ● Mercury—197 (¹⁹⁷ Hg) | 9 | D | 65.00 |
| 3 | Key in two of the following | | | |
| | three quantities: | | | |
| | ● Activity at time zero | A ₀ | A | A ₀ |
| | ● Time elapsed in days.hours | | | |
| | format* | t (dd.hh) | B | t (dd.hh) |
| | ● Present activity | A | C | A |
| 4 | Compute remaining variable: | | | |
| | ● Activity at time zero | | A | A ₀ |
| | ● Time elapsed in days.hours | | | |
| | format | | B | t (dd.hh) |
| | ● Present activity | | C | A |

| STEP | INSTRUCTIONS | INPUT DATA/UNITS | KEYS | OUTPUT DATA/UNITS |
|------|----------------------------------|---------------------|------|----------------------|
| 5 | For a new isotope, go to step 2; | | | |
| | to change one or both input | | | |
| | parameters, go to step 3. | | | |
| | | | | |
| | *Recall that two digits must | | | |
| | always be allocated for hours. | | | |
| | For example, 1 day 6 hours is | | | |
| | keyed in as 1.06. | | | |

Example:

An activity of 200 μCi is measured for a standard of ^{51}Cr . What is the activity after a week?

Keystrokes:

f **A** _____ \rightarrow
200 **A** _____ \rightarrow
7 **B** _____ \rightarrow
C _____ \rightarrow

Outputs:

667.20 ($\tau_{1/2}$ for ^{51}Cr)
200.00 (A_0)
7.00 ($t = 7$ days)
167.97 (A , μCi)

RADIOIMMUNOASSAY



This program performs the calculations for a logit/log plot of radioimmunoassay data. The program allows for any number of replicates in the counts input and for any number of standards. Outputs include correlation coefficient r , slope m , and intercept b of the least-squares regression line computed. Then, given counts for an unknown, the program will compute the corresponding concentration.

To run this program, first press **A** to initialize. Then key in the non-specific binding (or blank) counts, NSB, and press **B**; repeat for as many replicates as desired. After all replicates have been keyed in, press **R/S** to compute the average non-specific binding count. (This step is *not* optional; do not omit it.) The same procedure is repeated for the counts at zero concentration, B_0 , which are input to key **C**. After input of all replicates **R/S** is pressed to compute the average B_0 .

The next step in the operation of the program is the input of the data for the standards. The counts for the first standard are input to key **D**; as many replicates as desired may be keyed in. After all replicates for the first standard have been keyed in, the concentration of the standard is input to key **E**. This procedure (replicates to key **D**, concentration to key **E**) is repeated for as many standards as desired. Pressing key **f A** will then cause the output of the correlation coefficient r , the slope m , and the intercept b of the least-squares logit/log regression line computed from the standards. (The values of r , m , and b must be found before going to the next step, which is the calculation of the concentration of an unknown.) The regression performed is an unweighted regression.

At this point, the counts of an unknown may be keyed into **f B**; repeat for any number of replicates. After all replicates have been keyed in, **f C** may be pressed to find the concentration of that unknown. Repeat for as many unknowns as desired.

Two output options are available in this program. If neither option is selected, the only values output will be r , m , b , and the concentration of each unknown. Selection of the PRINT mode on key **f D** allows output of the following values as well: all input values (counts and standard concentrations) and the average of each set of counts input (assuming replicates). The second option, on key **f E**, is called PLOT. If this option is selected, the net B/B_0 and the log and logit (x and y) values for standards and unknowns will also be output. This information is intended to assist those who wish to make a plot by hand of the logit-log relationship.

Equations:

Let

NSB = average of replicate counts for non-specific binding

B_0 = average of replicate counts for zero concentration

B_i = average of replicate counts for i^{th} standard ($i = 1, 2, \dots, n$)

C_i = concentration of i^{th} standard

Let

$$x_i = \log C_i$$

$$\begin{aligned} y_i &= \text{logit} \left(\frac{B_i - \text{NSB}}{B_0 - \text{NSB}} \right) \\ &= \ln \left[\frac{(B_i - \text{NSB})/(B_0 - \text{NSB})}{1 - (B_i - \text{NSB})/(B_0 - \text{NSB})} \right] \\ &= \ln \left(\frac{B_i - \text{NSB}}{B_0 - B_i} \right) \end{aligned}$$

$$\text{net } B_i/B_0 = \frac{B_i - \text{NSB}}{B_0 - \text{NSB}}$$

The program fits a line of the form $y = mx + b$ to the (x_i, y_i) pairs. All sums below are from 1 to n .

$$m = \frac{\sum xy - \frac{\sum x \sum y}{n}}{\sum x^2 - \frac{(\sum x)^2}{n}}$$

$$b = \bar{y} - m \bar{x}$$

where:

$$\bar{y} = \frac{\sum y}{n}$$

$$\bar{x} = \frac{\sum x}{n}$$

$$r = \frac{\sum xy - \frac{\sum x \sum y}{n}}{\left[\sum x^2 - \frac{(\sum x)^2}{n} \right]^{1/2} \left[\sum y^2 - \frac{(\sum y)^2}{n} \right]^{1/2}}$$

Let

B = average of replicate counts for an unknown

C_u = concentration of unknown

$C_u = 10^x$

where $x = \frac{1}{m} \left[\ln \left(\frac{B - NSB}{B_0 - B} \right) - b \right]$

Remarks:



- 1. The term “intercept” is used in this program to refer to the point on the logit axis (the y-axis) where it is intersected by the regression line. It does not mean, as it is sometimes used in RIA documents, the concentration for which the value of the logit function is zero.
- 2. After computation of r, m, and b, these values may be found in the following registers: r in R_C and Z, m in R_B and Y, and b in R_A and X.

References:

Rodbard, Bridson, and Rayford, “Rapid calculation of radioimmunoassay results”, *J. Lab. Clin. Med.*, 74:770 (1969).

| STEP | INSTRUCTIONS | INPUT DATA/UNITS | KEYS | OUTPUT DATA/UNITS |
|------|-----------------------------------|------------------|-------------------|-------------------|
| 1 | Load side 1 and side 2. | | | |
| 2 | To allow output of input data | | | |
| | and intermediate results, turn | | | |
| | print function on. | | f D | 1.00 |
| 3 | To turn print function off later. | | f D | 0.00 |
| 4 | To allow output of (log conc., | | | |
| | logit) values, turn plot | | | |
| | function on. | | f E | 1.00 |
| 5 | To suppress further output of | | | |
| | plot data. | | f E | 0.00 |
| | Setup | | | |
| 6 | Initialize. | | A | |
| 7 | Key in non-specific binding | | | |
| | counts; repeat for as many | | | |
| | replicates as desired. | NSB | B | i |
| 8 | After all replicates, find | | | |
| | average NSB. | | R/S | <u>NSB</u> |

| STEP | INSTRUCTIONS | INPUT DATA/UNITS | KEYS | OUTPUT DATA/UNITS |
|------|---------------------------------------|------------------|------------|-------------------|
| 9 | Key in counts for zero dose; | | | |
| | repeat for as many replicates | | | |
| | as desired. | B_0 | C | i |
| 10 | After all replicates, find | | | |
| | average B_0 . | | R/S | \bar{B}_0 |
| | Standards | | | |
| 11 | Key in counts for first standard; | | | |
| | repeat for as many replicates | | | |
| | as desired. | B | D | i |
| 12 | Key in concentration of first | | | |
| | standard; optional outputs | | | |
| | are shown in parentheses; | | | |
| | 1.00 indicates first standard. | Conc. | E | (\bar{B}) |
| | | | | (net B/B_0) |
| | | | | (Conc.) |
| | | | | (Logit) |
| | | | | (Log conc.) |
| | | | | 1.00 |
| 13 | Repeat steps 11 and 12 for | | | |
| | all standards. | | | |
| | Results | | | |
| 14 | Calculate correlation coef- | | | |
| | ficient (r), slope (m), and | | | |
| | intercept (b) of regression line. | | f A | r |
| | | | | m |
| | | | | b |
| | Unknowns | | | |
| 15 | Key in counts for an unknown; | | | |
| | repeat for as many replicates | | | |
| | as desired. | B | f B | i |

| STEP | INSTRUCTIONS | INPUT DATA/UNITS | KEYS | OUTPUT DATA/UNITS |
|------|--------------------------------|---------------------|---|----------------------|
| 16 | Find concentration of | | | |
| | unknown; optional outputs | | | |
| | are shown in parentheses. | |   | (\bar{B}) |
| | | | | (net B/B_0) |
| | | | | Conc. |
| | | | | (Logit) |
| | | | | (Log) |
| 17 | Repeat steps 15 and 16 for | | | |
| | any number of unknowns. | | | |
| | New Case | | | |
| 18 | For a new assay, go to step 6. | | | |

Example:

Below are the data for non-specific binding (NSB), zero concentration (B_0), and various standards for a radioimmunoassay.

| Description | Counts per minute | Concentration (pg) |
|-------------|---------------------|--------------------|
| NSB | 425, 339, 342, 369 | - |
| B_0 | 10670, 10570, 10925 | - |
| Standard 1 | 9176, 9850 | 25 |
| Standard 2 | 8453, 7967 | 50 |
| Standard 3 | 6323, 6057 | 100 |
| Standard 4 | 3866, 4088 | 200 |
| Standard 5 | 2027, 2221 | 400 |
| Standard 6 | 1251, 1462 | 800 |

Find r , m , and b for the regression line. Find the concentrations corresponding to the unknown counts below.

| Unknown | Counts per minute |
|---------|-------------------|
| 1 | 10230, 10170 |
| 2 | 3270, 3400 |

Use the PRINT and PLOT options for complete outputs.

Keystrokes:

Outputs:

| | |
|------------------------------|--|
| A f D → | 1.00 (Print on) |
| f E → | 1.00 (Plot on) |
| 425 B → | 425.00 *** (1 st NSB) |
| | 1.00 |
| 339 B → | 339.00 *** |
| | 2.00 |
| 342 B → | 342.00 *** |
| | 3.00 |
| 369 B → | 369.00 *** |
| | 4.00 |
| R/S → | 368.75 *** (Avg. NSB) |
| 10670 C → | 10670.00 *** (1 st B ₀) |
| | 1.00 |
| 10570 C → | 10570.00 *** |
| | 2.00 |
| 10925 C → | 10925.00 *** |
| | 3.00 |
| R/S → | 10721.67 *** (Avg. B ₀) |
| 9176 D → | 9176.00 *** (1 st of std. 1) |
| | 1.00 |
| 9850 D → | 9850.00 *** |
| | 2.00 |
| 25 E → | 9513.00 *** (Avg. for std. 1) |
| | 0.88 *** (net B ₁ /B ₀) |
| | 25.00 *** (Conc. of std. 1) |
| | 2.02 *** (Logit = y ₁) |
| | 1.40 *** (Log = x ₁) |
| | 1.00 (Std. 1) |
| 8453 D → | 8453.00 *** (1 st of std. 2) |
| | 1.00 |

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| | |
|-----------------|---|
| 7967 D → | 7967.00 *** 2.00 |
| 50 E → | 8210.00 *** (Avg. for std. 2) 0.76 *** (net B ₂ /B ₀) 50.00 *** (Conc. of std. 2) 1.14 *** (y ₂) 1.70 *** (x ₂) 2.00 (Std. 2) |
| 6323 D → | 6323.00 *** (1 st of std. 3) 1.00 |
| 6057 D → | 6057.00 *** 2.00 |
| 100 E → | 6190.00 *** (Avg. for std. 3) 0.56 *** (net B ₃ /B ₀) 100.00 *** (Conc. of std. 3) 0.25 *** (y ₃) 2.00 *** (x ₃) 3.00 (Std. 3) |
| 3866 D → | 3866.00 *** (1 st of std. 4) 1.00 |
| 4088 D → | 4088.00 *** 2.00 |
| 200 E → | 3977.00 *** (Avg. for std. 4) 0.35 *** (net B ₄ /B ₀) 200.00 *** (Conc. of std. 4) -0.63 *** (y ₄) 2.30 *** (x ₄) 4.00 (Std. 4) |
| 2027 D → | 2027.00 *** (1 st of std. 5) 1.00 |
| 2221 D → | 2221.00 *** 2.00 |

| | |
|---------------------------|--|
| 400 E → | 2124.00 *** (Avg. for std. 5) 0.17 *** (net B_5/B_0) 400.00 *** (Conc. of std. 5) -1.59 *** (y_5) 2.60 *** (x_5) 5.00 (Std. 5) |
| 1251 D → | 1251.00 *** (1 st of std. 6) 1.00 |
| 1462 D → | 1462.00 *** 2.00 |
| 800 E → | 1356.50 *** (Avg. for std. 6) 0.10 *** (net B_6/B_0) 800.00 *** (Conc. of std. 6) -2.25 *** (y_6) 2.90 *** (x_6) 6.00 (Std. 6) |
| f A → | -1.00 *** (r) -2.89 *** (Slope m) 6.03 *** (Intercept b) |
| 10230 f B → | 10230.00 *** (1 st of unkn. 1) 1.00 |
| 10170 f B → | 10170.00 *** 2.00 |
| f C → | 10200.00 *** (Avg. of unkn. 1) 0.95 *** (net B/B_0) 11.83 *** (Conc. of unkn. 1) 2.94 *** (Unkn. y) 1.07 *** (Unkn. x) 11.83 |
| 3270 f B → | 3270.00 *** (1 st of unkn. 2) 1.00 |

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3400 **f** **B** →

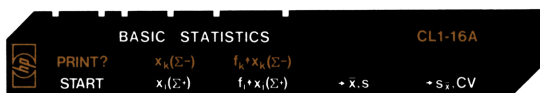
3400.00 ***
2.00

f **C** →

3335.00 *** (Avg. of unkn. 2)
0.29 *** (net B/B₀)
254.57 *** (Conc. of unkn. 2)
-0.91 *** (Unkn. y)
2.41 *** (Unkn. x)
254.57

Notes

BASIC STATISTICS



This program computes the basic statistics of one variable: mean (\bar{x}), standard deviation (s), standard error ($s_{\bar{x}}$), and coefficient of variation (C.V. %).

The input data to the program may be either grouped or ungrouped. Ungrouped data should be input to key **B** and grouped data to key **C**; keys **f B** and **f C** provide error correction for the ungrouped and grouped cases, respectively. If an incorrect entry is made, it may be corrected by keying in that entry a second time and pressing the appropriate error correction key. Suppose, for example, that 7.31 is one data point in a set of ungrouped data, but that a mistake is made in entering it. Instead of 7.31, the value 4.31 is input to key **B**. To correct this mistake, you would simply key in 4.31 and press **f B**. At this point the error has been eliminated. Now enter the correct data, 7.31, and press **B**.

Equations:

Ungrouped data:

Let $\{x_1, x_2, \dots, x_n\}$ be the set of data points.

$$\text{Mean } \bar{x} = \frac{1}{n} \sum_{i=1}^n x_i$$

$$s = \sqrt{\frac{\sum x_i^2 - \frac{(\sum x_i)^2}{n}}{n - 1}}$$

$$\text{Standard error } s_{\bar{x}} = \frac{s}{\sqrt{n}}$$

$$\text{Coefficient of variation C.V. \%} = \frac{s}{\bar{x}} \times 100$$

Grouped data:

Let $\{x_1, x_2, \dots, x_n\}$ be a set of data points occurring with the respective frequencies f_1, f_2, \dots, f_n .

$$\text{Mean } \bar{x} = \frac{\sum f_i x_i}{\sum f_i}$$

$$\text{Standard deviation } s = \sqrt{\frac{\sum f_i x_i^2 - \frac{(\sum f_i x_i)^2}{\sum f_i}}{\sum f_i - 1}}$$

$$\text{Standard error } s_{\bar{x}} = \frac{s}{\sqrt{\sum f_i}}$$

$$\text{Coefficient of variation C. V. \%} = \frac{s}{\bar{x}} \times 100$$

Remarks:

1. Grouped and ungrouped data may be mixed in the same set of data.
2. The preprogrammed $\Sigma+$ and $\Sigma-$ keys may be used to input and correct ungrouped data in place of keys **B** and **f B**. Calculation of mean and standard deviation may also be done by the preprogrammed keys \bar{x} and **s** for both grouped and ungrouped data.

| STEP | INSTRUCTIONS | INPUT DATA/UNITS | KEYS | OUTPUT DATA/UNITS |
|------|---|------------------|--------------|-------------------|
| 1 | Load side 1. | | | |
| 2 | Initialize. | | A | 0.00 |
| 3 | To allow printing of input data, turn print function on. | | f A | 1.00 |
| 4 | To turn print function off later. | | f A | 0.00 |
| 5 | For ungrouped data, go to step 6; for grouped data, go to step 9. | | | |
| | Ungrouped data | | | |
| 6 | Perform this step for $i = 1,$ $2, \dots, n:$ | | | |
| | Input data point. | x_i | B | i |
| 7 | To correct an erroneous entry. | x_k | f B | i |
| 8 | Go to step 11. | | | |
| | Grouped data | | | |
| 9 | Perform this step for $i = 1,$ $2, \dots, n:$ | | | |
| | Input frequency and data. | f_i | ENTER | |
| | | x_i | C | i |

| STEP | INSTRUCTIONS | INPUT DATA/UNITS | KEYS | OUTPUT DATA/UNITS |
|------|--|------------------|--------------|-------------------|
| 10 | To correct an erroneous entry. | f_k | ENTER | |
| | | x_k | f C | i |
| | Results | | | |
| 11 | Compute mean and standard deviation. | | D | \bar{x} |
| | | | | s |
| 12 | Compute standard error and coefficient of variation. | | E | $s_{\bar{x}}$ |
| | | | | C.V. % |
| 13 | For a new set of data, go to step 2. | | | |

Example 1:

Hemoglobin concentration was measured for nine male patients. Compute the basic statistics for these data.

Hemoglobin concentration (g/dl)

| | |
|------|------|
| 13.8 | 17.4 |
| 16.9 | 13.4 |
| 16.5 | 17.9 |
| 17.7 | 15.2 |
| 16.0 | |

Keystrokes:

| | |
|---------------|---|
| A | → |
| f A | → |
| 13.8 B | → |
| 16.9 B | → |
| 16.5 B | → |
| 17.7 B | → |

Outputs:

| |
|---------------------|
| 0.00 |
| 1.00 (Print on) |
| 13.80 *** (x_i) |
| 1.00 (i) |
| 16.90 *** |
| 2.00 |
| 16.50 *** |
| 3.00 |
| 17.70 *** |
| 4.00 |

| | | | |
|------|-----|---|----------------------------|
| 16 | B | → | 16.00 *** |
| | | | 5.00 |
| 17.4 | B | → | 17.40 *** |
| | | | 6.00 |
| 3.4 | B | → | 3.40 *** (Error!) |
| | | | 7.00 (k = 7) |
| 3.4 | f B | → | 3.40 *** (Correction) |
| | | | 6.00 |
| 13.4 | B | → | 13.40 *** (x_7) |
| | | | 7.00 |
| 17.9 | B | → | 17.90 *** |
| | | | 8.00 |
| 15.2 | B | → | 15.20 *** |
| | | | 9.00 |
| D | | → | 16.09 *** (Mean) |
| | | | 1.65 *** (Std. dev.) |
| E | | → | 0.55 *** ($S_{\bar{x}}$) |
| | | | 10.23 *** (C.V. %) |

Example 2:

A certain test was performed on college students ranging in age from 18 to 22 years. The number of subjects of each age is shown in the table. Compute the mean age of the students in the test.

| | | | | | |
|------------|----|----|----|----|----|
| Age | 18 | 19 | 20 | 21 | 22 |
| # Subjects | 5 | 9 | 13 | 7 | 1 |


Keystrokes:

| | |
|--|---|
| A | → |
| If Example 1 has just been run, turn print off: | |
| f A | → |
| 5 ENTER 18 C | → |
| 9 ENTER 19 C | → |
| 13 ENTER 20 C | → |
| 7 ENTER 21 C | → |
| 1 ENTER 22 C | → |
| D | → |

Outputs:

| |
|----------------------|
| 0.00 |
| |
| 0.00 (Print off) |
| 1.00 |
| 2.00 |
| 3.00 |
| 4.00 |
| 5.00 |
| 19.71 *** (Mean) |
| 1.05 *** (Std. dev.) |

CHI-SQUARE EVALUATION AND DISTRIBUTION

| | CHI-SQUARE | TEST | AND | DISTRIBUTION | CL1-17A |
|---|------------|-----------------|---------------------------|-------------------------|---------------------------|
|  | PRINT? | $O_k(\Sigma^-)$ | $O_k \cdot E_k(\Sigma^-)$ | $x \cdot f(x)$ | $x \cdot P(x)$ |
| | START | $O, (\Sigma^+)$ | $O \cdot E, (\Sigma^+)$ | $\rightarrow \chi^2; E$ | $\nu \cdot \Gamma(\nu/2)$ |

This program allows you to perform two important calculations concerning the chi-square statistic. The first of these calculates the value of the χ^2 statistic for the goodness of fit test. The second evaluates the chi-square density $f(x)$ and the cumulative distribution $P(x)$ given x and the degrees of freedom ν .

The χ^2 statistic may be computed for the case where the expected frequencies are equal as well as for the case where they are different. If they are equal, only the observed frequencies O_i need be input to key **B**; error correction is available on key **f B**. After calculation of χ^2 from key **D**, the expected frequency E may be calculated. If the expected frequencies are different, both the observed and expected frequencies should be input to key **C**. Error correction is provided on key **f C**.

To make calculations involving the chi-square distribution, first input the degrees of freedom ν to key **E**. Then key in the value of x and press **f D** to find the density $f(x)$ or **f E** to find the cumulative distribution $P(x)$.

Equations:

Chi-square evaluation:

$$\chi^2 = \sum_{i=1}^n \frac{(O_i - E_i)^2}{E_i}$$

where:

O_i = observed frequency

E_i = expected frequency

If the expected values are equal

$$\left(E = E_i = \frac{\Sigma O_i}{n} \text{ for all } i \right)$$

then

$$\chi^2 = \frac{n \Sigma O_i^2}{\Sigma O_i} - \Sigma O_i$$

Chi-square distribution:

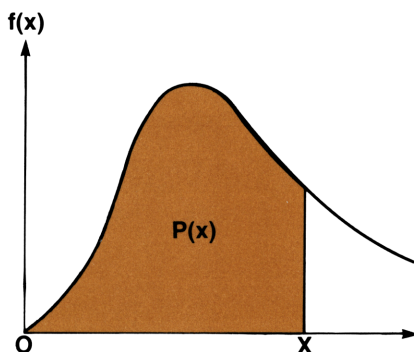
Chi-square density:

$$f(x) = \frac{1}{2^{\frac{\nu}{2}} \Gamma\left(\frac{\nu}{2}\right)} x^{\frac{\nu}{2}-1} e^{-\frac{x}{2}}$$

where:

$$x \geq 0$$

ν is the degrees of freedom.



Series approximation is used to evaluate the cumulative distribution

$$P(x) = \int_0^x f(t) dt$$

$$= \left(\frac{x}{2}\right)^{\frac{\nu}{2}} \frac{e^{-\frac{x}{2}}}{\Gamma\left(\frac{\nu+2}{2}\right)} \left[1 + \sum_{k=1}^{\infty} \frac{x^k}{(\nu+2)(\nu+4) \dots (\nu+2k)} \right]$$

where:

$$\Gamma\left(\frac{\nu}{2}\right) = \begin{cases} \left(\frac{\nu}{2} - 1\right)!, & \nu \text{ even} \\ \left(\frac{\nu}{2} - 1\right)\left(\frac{\nu}{2} - 2\right) \dots \left(\frac{1}{2}\right) \Gamma\left(\frac{1}{2}\right), & \nu \text{ odd} \end{cases}$$

$$\Gamma\left(\frac{1}{2}\right) = \sqrt{\pi}$$

The program computes successive partial sums of the above series. When two consecutive partial sums are equal, the value is used as the sum of the series.

Remarks:

- 1. In order to apply the goodness of fit test to a set of given data, it may be necessary to combine some classes to ensure that each expected frequency is not too small (not less than, say, 5).
- 2. The program for distribution requires that $\nu \leq 141$. If $\nu > 141$, erroneous overflow will result.
- 3. If both x and ν are large, the calculation of $f(x)$ may cause overflow.

References:

(Evaluation) J.E. Freund, *Mathematical Statistics*, Prentice Hall, 1962.
(Distribution) Abramowitz and Stegun, *Handbook of Mathematical Functions*, National Bureau of Standards, 1968.

| STEP | INSTRUCTIONS | INPUT DATA/UNITS | KEYS | OUTPUT DATA/UNITS |
|------|---|------------------|------|-------------------|
| 1 | Load side 1 and side 2. | | | |
| 2 | Initialize. | | A | 20.00 |
| 3 | To allow printing of data and results, turn the print function on. | | I A | 1.00 |
| 4 | To turn the print function off later. | | I A | 0.00 |
| 5 | For χ^2 evaluation, go to step 6; for χ^2 distribution, go to step 15. χ^2 evaluation | | | |
| 6 | If the expected frequencies are equal, go to step 7; if they are not equal, go to step 11. Expected frequencies equal | | | |
| 7 | Perform this step for $i = 1, 2, \dots, n$: Key in observed value. | O_i | B | i |

| STEP | INSTRUCTIONS | INPUT DATA/UNITS | KEYS | OUTPUT DATA/UNITS |
|------|---|------------------|----------------------------|-------------------|
| 8 | To correct an erroneous entry. | O_k | f B | i |
| 9 | Calculate the χ^2 statistic and (optionally) the average expected frequency. | | D | χ^2 |
| | | | R/S | E |
| 10 | For a new case, go to step 2. | | | |
| | Expected frequencies | | | |
| | unequal | | | |
| 11 | Perform this step for $i = 1,$ 2, ..., n: | | | |
| | Key in observed and expected frequency. | O_i E_i | ENTER C | i |
| 12 | To correct an erroneous entry. | O_k E_k | ENTER f C | i |
| 13 | Calculate the χ^2 statistic. | | D | χ^2 |
| 14 | For a new case, go to step 2. | | | |
| | χ^2 distribution | | | |
| 15 | Key in degrees of freedom. | ν | E | $\Gamma(\nu/2)$ |
| 16 | Key in x and compute either | | | |
| | • Density | x | f D | f(x) |
| | or | | | |
| | • Cumulative distribution | x | f E | P(x) |
| 17 | For a new case, go to step 2. | | | |

Example:

Ten one-minute counts of a Cesium-137 check source yielded the following results. Use this program to evaluate the counting instrument. (Note that with 10 data points, the degrees of freedom $\nu = 9$.)

| | |
|-------|-------|
| 25601 | 25553 |
| 25546 | 25841 |
| 25592 | 25560 |
| 25820 | 25633 |
| 25569 | 25464 |

Keystrokes:

A **f** **A** →
25601 **B** →
25546 **B** →
25592 **B** →
25820 **B** →
25569 **B** →
25553 **B** →
25841 **B** →
25560 **B** →
25633 **B** →
25464 **B** →
D →
R/S →
9 **E** →
5.10 **f** **E** →

Outputs:

1.00 (Print on)
25601.00 ***
1.00
25546.00 ***
2.00
25592.00 ***
3.00
25820.00 ***
4.00
25569.00 ***
5.00
25553.00 ***
6.00
25841.00 ***
7.00
25560.00 ***
8.00
25633.00 ***
9.00
25464.00 ***
10.00
5.10 *** (χ^2)
25617.90 *** (E)
9.00 *** (ν)
11.63 *** ($\Gamma(\nu/2)$)
5.10 ***
0.17 *** (P (χ^2))

Since P (χ^2) is between 0.1 and 0.9, the counting instrument is assumed to be operating properly.

Notes

t STATISTICS

| | | | | |
|--------|------------------------|-----------------------------|-----------------------|---------------------|
| | t STATISTICS | | CL1-18A | |
| PRINT? | $x_k + y_k (\Sigma -)$ | | $x_k, y_k (\Sigma -)$ | d |
| START | $x_i + y_i (\Sigma +)$ | $\rightarrow t: df, D, S_D$ | $x_i, y_i (\Sigma +)$ | $\rightarrow t: df$ |

This program will compute either of two test statistics which are used to compare population means: the paired t statistic or the t statistic for two means.

The paired t statistic applies to a set of *paired* observations drawn from two normal populations with unknown means μ_1, μ_2 :

| | | | | |
|-------|-------|-------|-----|-------|
| x_i | x_1 | x_2 | ... | x_n |
| y_i | y_1 | y_2 | ... | y_n |

The paired t statistic can be used to test the validity of the hypothesis that the means are equal. If the computed value of t is significant (as determined by *t Distribution*, CL1-19A), then we reject the hypothesis that the population means are equal.

The x- and y-values are input to key **B**. Error correction is provided by key **f B**. After the input of all x-y pairs, the t statistic may be found by pressing **C**.

The t statistic for two means applies to independent random samples $\{x_1, x_2, \dots, x_{n_1}\}$ and $\{y_1, y_2, \dots, y_{n_2}\}$ drawn from two normal populations with unknown means μ_1, μ_2 and the same unknown variance σ^2 . The t statistic is used to test the validity of the hypothesis that the populations means differ by some amount d (i.e., that $\mu_1 - \mu_2 = d$). Note that d may be chosen to be zero.

To operate this routine, the x-values should first be keyed in to key **D**. Error correction is available on key **f D**. After all x-values have been input, the value of d should be input to key **f E**. Then the y-values should be keyed in to key **D**. After input of all the y-values, the t statistic may be found by pressing **E**.

Equations:*Paired t statistic*

let

$$D_i = x_i - y_i$$

$$\bar{D} = \frac{1}{n} \sum_{i=1}^n D_i$$

$$s_D = \sqrt{\frac{\sum D_i^2 - \frac{1}{n} (\sum D_i)^2}{n - 1}}$$

$$s_{\bar{D}} = \frac{s_D}{\sqrt{n}}$$

The test statistic

$$t = \frac{\bar{D}}{s_{\bar{D}}}$$

which has $n - 1$ degrees of freedom (df) can be used to test the null hypothesis

$$H_0: \mu_1 = \mu_2$$

t statistic for two means

Define

$$\bar{x} = \frac{1}{n_1} \sum_{i=1}^{n_1} x_i$$

$$\bar{y} = \frac{1}{n_2} \sum_{i=1}^{n_2} y_i$$

$$t = \frac{\bar{x} - \bar{y} - d}{\sqrt{\frac{1}{n_1} + \frac{1}{n_2}} \sqrt{\frac{\sum x_i^2 - n_1 \bar{x}^2 + \sum y_i^2 - n_2 \bar{y}^2}{n_1 + n_2 - 2}}}$$

We can use this t statistic which has the t distribution with $n_1 + n_2 - 2$ degrees of freedom (df) to test the null hypothesis

$$H_0: \mu_1 - \mu_2 = d$$

References:

(Paired t) B. Ostle, *Statistics in Research*, Iowa State University Press, 1963.

(t for two means) K.A. Brownlee, *Statistical Theory and Methodology in Science and Engineering*, John Wiley and Sons, 1965.

| STEP | INSTRUCTIONS | INPUT DATA/UNITS | KEYS | OUTPUT DATA/UNITS |
|------|--|---------------------|---------------|----------------------|
| 1 | Load side 1 and 2 of program. | | | |
| 2 | Initialize. | | A | |
| 3 | To allow output of data and results, turn print function on. | | f A | 1.00 |
| 4 | To turn print function off later. | | f A | 0.00 |
| 5 | For t statistic for two means, go to step 11; for paired t statistic, go to step 6. | | | |
| | Paired t statistic | | | |
| 6 | Repeat this step for all data pairs ($i = 1, 2, \dots, n$): | | | |
| | Key in x- and y-values. | x_i | ENTER* | |
| | | y_i | B | i |
| 7 | To correct an erroneous entry. | x_k | ENTER* | |
| | | y_k | f B | i |
| 8 | Compute paired t statistic. | | C | t |
| 9 | (optional) Compute degrees of freedom, mean difference, and standard deviation of D. | | | |
| | | | R/S | df |
| | | | | \bar{D} . |
| | | | | s_D |
| 10 | For a new case, go to step 2. | | | |
| | t statistic for two means | | | |
| 11 | Repeat this step for all x-values ($i = 1, 2, \dots, n_1$): | | | |
| | Key in x-value. | x_i | D | i |
| 12 | To correct an erroneous entry. | x_k | f D | i |

| STEP | INSTRUCTIONS | INPUT DATA/UNITS | KEYS | OUTPUT DATA/UNITS |
|------|--|------------------|---------------------|-------------------|
| 13 | Key in difference to be tested. | d | f E | d |
| 14 | Repeat this step for all y-values ($i = 1, 2, \dots, n_2$): | | | |
| | Key in y-value. | y_i | D | i |
| 15 | To correct an erroneous entry. | y_k | f D | i |
| 16 | Compute t statistic for two means. | | E | t |
| 17 | (optional) Compute degrees of freedom. | | R/S | df |
| 18 | (optional) Change value of d and repeat step 16. | d | STO 7 | |
| 19 | For a new case go to step 2. | | | |

Example 1:

The hemoglobin concentration in blood samples from six patients was measured by two different methods. Use the paired t-statistic to determine if there is a significant difference between the two methods of measurement.

| Sample | Method | |
|--------|----------|----------|
| | 1 (g/dl) | 2 (g/dl) |
| 1 | 17.6 | 17.4 |
| 2 | 13.0 | 12.9 |
| 3 | 15.3 | 15.3 |
| 4 | 15.0 | 15.2 |
| 5 | 15.0 | 15.0 |
| 6 | 14.6 | 14.5 |

Keystrokes:

A \longrightarrow
f **A** \longrightarrow
 17.6 **ENTER** 17.4 **B** \longrightarrow

Outputs:

0.00
 1.00 (Print on)
 17.60 *** (x_1)
 17.40 *** (y_1)
 1.00 ($i = 1$)

13 **ENTER** 12.9 **B** →

15.3 **ENTER** 15.2 **B** →

15.3 **ENTER** 15.2 **f** **B** →

15.3 **ENTER** 15.3 **B** →

15 **ENTER** 15.2 **B** →

15 **ENTER** 15 **B** →

14.6 **ENTER** 14.5 **B** →

C →

R/S →

13.00 ***
12.90 ***
2.00
15.30 ***
15.20 *** (Error!)
3.00
15.30 ***
15.20 *** (Corrected)
2.00
15.30 ***
15.30 ***
3.00
15.00 ***
15.20 ***
4.00
15.00 ***
15.00 ***
5.00
14.60 ***
14.50 ***
6.00
0.60 *** (t)
5.00 *** (df)
0.03 *** (\bar{D})
0.14 *** (S_D)

To interpret these results, load *t Distribution* (CL1-19A) and find the cumulative distribution I(x) for $x = 0.60$ and 5 degrees of freedom.

Keystrokes:

5 **A** .60 **D** →

Outputs:

0.43 *** (I (0.60))

The probability of $|t| > 0.60$ is thus 57%. We conclude that the hypothesis that the means are equal cannot be rejected.

Example 2:

Hemoglobin concentration was measured for nine male and seven female patients. Use the t-statistic for two means to test the hypothesis that the difference between the means is negligible (i.e., $d = 0$).

| <u>Hgb concentration (g/dl)</u> | |
|---------------------------------|--------------|
| <u>Men</u> | <u>Women</u> |
| 13.8 | 11.9 |
| 16.9 | 14.4 |
| 16.5 | 13.7 |
| 17.7 | 16.8 |
| 16.0 | 11.7 |
| 17.4 | 14.9 |
| 13.4 | 12.3 |
| 17.9 | |
| 15.2 | |

Keystrokes:

Outputs:

| | | |
|-------------------------------------|-----------|-------------|
| A → | 0.00 | |
| If example 1 has not just been run: | | |
| f A → | 1.00 | (Print on) |
| 13.8 D → | 13.80 *** | (x_1) |
| 16.9 D → | 1.00 | ($i = 1$) |
| 16.5 D → | 16.90 *** | |
| 17.7 D → | 2.00 | |
| 16 D → | 16.50 *** | |
| 17.4 D → | 3.00 | |
| 13.4 D → | 17.70 *** | |
| 17.9 D → | 4.00 | |
| 15.2 D → | 16.00 *** | |
| 0 f E → | 5.00 | |
| | 17.40 *** | |
| | 6.00 | |
| | 13.40 *** | |
| | 7.00 | |
| | 17.90 *** | |
| | 8.00 | |
| | 15.20 *** | |
| | 9.00 | |
| | 0.00 *** | ($d = 0$) |

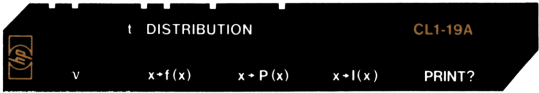
18-07

| | | | |
|----------------------------------|---|----------|---------------------|
| 11.9 | D | → | 11.90 *** (y_1) |
| | | | 1.00 ($i = 1$) |
| 14.4 | D | → | 14.40 *** |
| | | | 2.00 |
| 13.7 | D | → | 13.70 *** |
| | | | 3.00 |
| 16.8 | D | → | 16.80 *** |
| | | | 4.00 |
| 11.7 | D | → | 11.70 *** |
| | | | 5.00 |
| 14.9 | D | → | 14.90 *** |
| | | | 6.00 |
| 12.3 | D | → | 12.30 *** |
| | | | 7.00 |
| E | | → | 2.76 *** (t) |
| R/S | | → | 14.00 *** (df) |
| Load t Distribution (CL1-19A). | | | |
| 14 | A | 2.76 D → | 0.98 *** (I (2.76)) |

Thus the value of t is significant and we should reject the hypothesis that the average hemoglobin concentrations in males and females are equal.

Notes

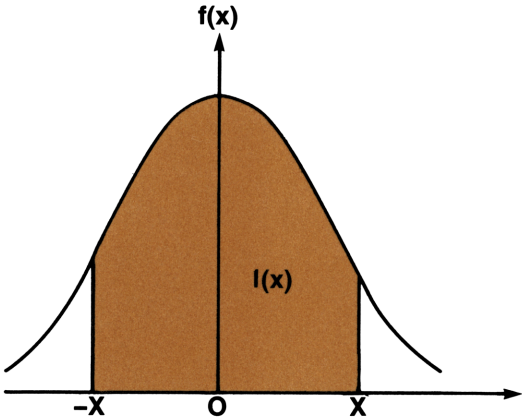
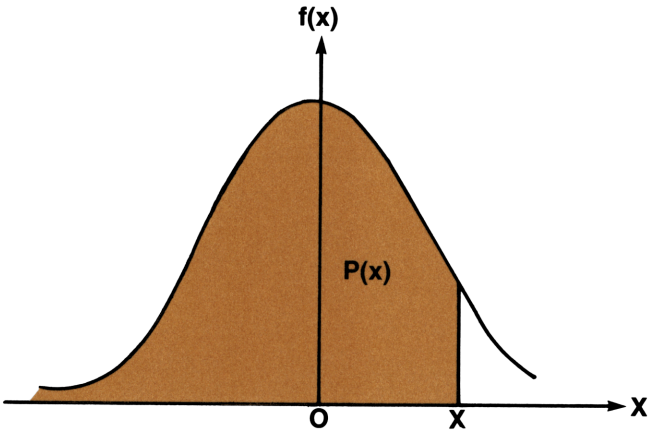
t DISTRIBUTION



This program calculates three parameters of the t distribution given x and the degrees of freedom ν . The density function $f(x)$ is computed as well as two measures of the area under the distribution curve, $P(x)$ and, for $x > 0$, $I(x)$, where

$$P(x) = \int_{-\infty}^x f(y) \, dy$$

and $I(x) = \int_{-x}^x f(y) \, dy$.



Equations:

$$f(x) = \frac{\Gamma\left(\frac{\nu+1}{2}\right)}{\sqrt{\pi\nu} \Gamma\left(\frac{\nu}{2}\right)} \left(1 + \frac{x^2}{\nu}\right)^{-\frac{\nu+1}{2}}$$

(1) ν even

$$I(x) = \sin \theta \left\{ 1 + \frac{1}{2} \cos^2 \theta + \frac{1 \cdot 3}{2 \cdot 4} \cos^4 \theta + \dots \right. \\ \left. + \frac{1 \cdot 3 \cdot 5 \dots (\nu-3)}{2 \cdot 4 \cdot 6 \dots (\nu-2)} \cos^{\nu-2} \theta \right\}$$

(2) ν odd

$$I(x) = \begin{cases} \frac{2\theta}{\pi} \text{ if } \nu = 1 \\ \frac{2\theta}{\pi} + \frac{2}{\pi} \cos \theta \left\{ \sin \theta \left[1 + \frac{2}{3} \cos^2 \theta + \dots \right. \right. \\ \left. \left. + \frac{2 \cdot 4 \dots (\nu-3)}{1 \cdot 3 \dots (\nu-2)} \cos^{\nu-3} \theta \right] \right\} \text{ if } \nu > 1 \end{cases}$$

where

$$\theta = \tan^{-1} \left(\frac{x}{\sqrt{\nu}} \right)$$

$$P(x) = \begin{cases} \frac{1 + I(x)}{2} \text{ if } x > 0 \\ \frac{1 - I(x)}{2} \text{ if } x \leq 0 \end{cases}$$

Remarks:

The program requires $\nu < 141$. Otherwise an erroneous overflow will result.

Reference:




Abramowitz and Stegun, *Handbook of Mathematical Functions*, National Bureau of Standards, 1970.

| STEP | INSTRUCTIONS | INPUT DATA/UNITS | KEYS | OUTPUT DATA/UNITS |
|------|----------------------------------|------------------|----------|-------------------|
| 1 | Load side 1 and side 2 | | | |
| | of program. | | | |
| 2 | To allow printing of inputs, | | | |
| | turn print function on. | | E | 1.00 |
| 3 | To turn print function off | | | |
| | later. | | E | 0.00 |
| 4 | Key in degrees of freedom. | ν | A | ν |
| 5 | Key in x and compute either | | | |
| | • Density function | x | B | f(x) |
| | or | | | |
| | • Cumulative distribution | x | C | P(x) |
| | or | | | |
| | • Integral, -x to x ($x > 0$). | x | D | I(x) |

Example 1:

Find the density function and P(x) for $x = 1.6$ with 9 degrees of freedom.

Keystrokes:

9 **A** 
1.6 **B** 
1.6 **C** 



Outputs:

9.00 (ν)
0.11 *** (f (x))
0.93 *** (P (x))

Example 2:

Find I(x) for $x = 1.83$ and $\nu = 11$.

Keystrokes:

11 **A** 
1.83 **D** 

Outputs:

11.00 (ν)
0.91 *** (I (x))

Notes

Notes

PROGRAM LISTINGS

The following listings are included for your reference. A table of keycodes and keystrokes corresponding to the symbols used in the listings can be found in Appendix E of your Owner's Handbook.

| Program | Page |
|--|---------------|
| 1. Beer's Law | L01-01 |
| 2. Protein Electrophoresis | L02-01 |
| 3. LDH Isoenzymes | L03-01 |
| 4. Body Surface Area | L04-01 |
| 5. Urea Clearance | L05-01 |
| 6. Creatinine Clearance | L06-01 |
| 7. Amniotic Fluid Assay | L07-01 |
| 8. Blood Acid-Base Status | L08-01 |
| 9. Oxygen Saturation and Content | L09-01 |
| 10. Red Cell Indices | L10-01 |
| 11. Total Blood Volume | L11-01 |
| 12. Schilling Test | L12-01 |
| 13. Thyroid Uptake | L13-01 |
| 14. Radioactive Decay Corrections | L14-01 |
| 15. Radioimmunoassay | L15-01 |
| 16. Basic Statistics | L16-01 |
| 17. Chi-square Evaluation and Distribution | L17-01 |
| 18. t Statistics | L18-01 |
| 19. t Distribution | L19-01 |

Beer's Law

| | | | | | | | | | |
|-----------|--|------------------|---|--------------|-----|----|------------------|------------------|------------------|
| 001 *LBL4 | A → %T. | 057 + | For % T _u , compute A _u . | | | | | | |
| 002 STOD | | 058 GTO0 | ----- | | | | | | |
| 003 F0? | | 059 *LBL4 | | | | | | | |
| 004 PRTX | A. | 060 CHS | | | | | | | |
| 005 2 | | 061 *LBL6 | | | | | | | |
| 006 - | | 062 ST09 | Store A _u . | | | | | | |
| 007 CHS | | 063 RCLW | Display input. | | | | | | |
| 008 I0? | | 064 RTN | ----- | | | | | | |
| 009 STOE | %T. | 065 *LBL6 | | | | | | | |
| 010 F0? | | 066 ST05 | | | | | | | |
| 011 PRTX | | 067 F0? | C _s → C _u . | | | | | | |
| 012 F0? | | 068 SPC | | | | | | | |
| 013 SFC | | 069 F0? | | | | | | | |
| 014 RTN | | 070 PRTX | | | | | | | |
| 015 *LBL5 | ----- | 071 RCL9 | | | | | | | |
| 016 STOE | %T → A. | 072 * | | | | | | | |
| 017 F0? | | 073 RCL6 | $C_u = C_s \times \frac{A_u}{A_s}$ | | | | | | |
| 018 PRTX | %T. | 074 ÷ | | | | | | | |
| 019 LOG | | 075 ST07 | | | | | | | |
| 020 CHS | | 076 F0? | | | | | | | |
| 021 2 | | 077 PRTX | | | | | | | |
| 022 + | | 078 F0? | | | | | | | |
| 023 STOD | | 079 SPC | | | | | | | |
| 024 F0? | | 080 RTN | ----- | | | | | | |
| 025 PRTX | A. | 081 *LBL6 | | | | | | | |
| 026 F0? | | 082 0 | Clear for reprint. | | | | | | |
| 027 SPC | | 083 ST04 | | | | | | | |
| 028 RTN | | 084 STOE | | | | | | | |
| 029 *LBL0 | + % T _s (-A _s). | 085 ST0C | | | | | | | |
| 030 ST0C | | 086 STOD | | | | | | | |
| 031 F0? | | 087 STOE | | | | | | | |
| 032 SPC | | 088 ST01 | | | | | | | |
| 033 F0? | | 089 RTN | | | | | | | |
| 034 PRTX | For absorbance, GTO 3. | 090 *LBL6 | Patient ID = Pnt# .01. | | | | | | |
| 035 X0? | | 091 INT | | | | | | | |
| 036 GTO3 | | 092 . | | | | | | | |
| 037 LOG | | 093 0 | | | | | | | |
| 038 CHS | | 094 1 | | | | | | | |
| 039 2 | For %T _s , compute A _s . | 095 + | | | | | | | |
| 040 + | ----- | 096 ST01 | | | | | | | |
| 041 GTO0 | | 097 PRTX | | | | | | | |
| 042 *LBL3 | | 098 SPC | | | | | | | |
| 043 CHS | | 099 RTN | | | | | | | |
| 044 *LBL0 | | 100 *LBL4 | | | | | | | |
| 045 STOB | Store A _s . | 101 F0? | Print toggle. | | | | | | |
| 046 RCLC | Display input. | 102 GTO0 | | | | | | | |
| 047 RTN | ----- | 103 SF0 | | | | | | | |
| 048 *LBL0 | + % T _u (-A _u). | 104 1 | | | | | | | |
| 049 ST0A | | 105 RTN | | | | | | | |
| 050 F0? | | 106 *LBL0 | | | | | | | |
| 051 PRTX | For absorbance, GTO 4. | 107 CF0 | | | | | | | |
| 052 X0? | | 108 0 | | | | | | | |
| 053 GTO4 | | 109 RTN | | | | | | | |
| 054 LOG | | 110 *LBL6 | ----- | | | | | | |
| 055 CHS | | 111 SFC | Reprint | | | | | | |
| 056 2 | | 112 SPC | | | | | | | |
| REGISTERS | | | | | | | | | |
| 0 | 1 | 2 | 3 | 4 | 5 | 6 | 7 C _u | 8 C _s | 9 A _u |
| S0 | S1 | S2 | S3 | S4 | S5 | S6 | S7 | S8 | S9 |
| A | Input to [D] | B A _s | C | Input to [C] | D A | E | % T | I | Pnt# .01 |

| | | | |
|----------|--|--|--|
| 113 RCLT | | | |
| 114 INT | | | |
| 115 . | | | |
| 116 0 | | | |
| 117 1 | | | |
| 118 + | | | |
| 119 PRTX | Patient ID. | | |
| 120 SPC | A. | | |
| 121 RCLC | | | |
| 122 PRTX | %T. | | |
| 123 RCLC | | | |
| 124 PRTX | | | |
| 125 SPC | | | |
| 126 SPC | | | |
| 127 RCLC | + % T _s (-A _s). | | |
| 128 PRTX | | | |
| 129 RCLC | + % T _u (-A _u). | | |
| 130 PRTX | | | |
| 131 SPC | | | |
| 132 RCLC | C _s . | | |
| 133 PRTX | | | |
| 134 RCLC | C _u . | | |
| 135 PRTX | | | |
| 136 RTN | ----- | | |

| LABELS | | | | | FLAGS | | SET STATUS | | |
|---------|--------|---------------------------------------|---------------------------------------|----------------------------------|---------|--|---|---|--|
| A A→%T | B %T→A | C +%T _s (-A _s) | D +%T _u (-A _u) | E C _s →C _u | 0 Print | FLAGS | TRIG | DISP | |
| a Clear | b | c Ptnt # | d P off? | e Reprint | 1 | ON OFF 0 <input checked="" type="checkbox"/> <input type="checkbox"/> | DEG <input checked="" type="checkbox"/> | FIX <input checked="" type="checkbox"/> | |
| 0 Used | 1 | 2 | 3 Store A _s | 4 Store A _u | 2 | 1 <input type="checkbox"/> <input checked="" type="checkbox"/> | GRAD <input type="checkbox"/> | SCI <input type="checkbox"/> | |
| 5 | 6 | 7 | 8 | 9 | 3 | 2 <input type="checkbox"/> <input checked="" type="checkbox"/> | RAD <input type="checkbox"/> | ENG <input type="checkbox"/> | |
| | | | | | | 3 <input type="checkbox"/> <input checked="" type="checkbox"/> | | n <u>2</u> | |

Protein Electrophoresis

| | | | | | | | | | |
|----------------------|--|----------------------|---|----------------------|----|----|----|---------|-----------------------|
| 001 #LBLA | Initialize. | 057 STO! | Point to Fract ₁ . | | | | | | |
| 002 2 | | 058 #LBLB | ----- | | | | | | |
| 003 5 | | 059 RCL i | | | | | | | |
| 004 STOI | | 060 RCL 0 | | | | | | | |
| 005 CLX | | 061 ÷ | | | | | | | |
| 006 STOI | | 062 RCL 2 | $\text{Gms} = \frac{\text{Fract}_i}{\Sigma} \times \text{T Pr}$ | | | | | | |
| 007 STOI | | 063 x | | | | | | | |
| 008 STOI | | 064 PRTY | | | | | | | |
| 009 STOI | | 065 RCL i | | | | | | | |
| 010 F0? | | 066 RCL 1 | | | | | | | |
| 011 SPC | | 067 X=Y? | Down to R _{25-n} ? | | | | | | |
| 012 RTN | ----- | 068 GTOP | Yes, exit. | | | | | | |
| 013 #LBLB | Input fractions. | 069 DSZ! | No, decrement and loop again. | | | | | | |
| 014 F0? | | 070 GTOP | ----- | | | | | | |
| 015 PRTX | Fract _i → R _{25-i} . | 071 #LBL0 | Display 0.00 and return. | | | | | | |
| 016 DSZ! | | 072 CLX | ----- | | | | | | |
| 017 STOI | | 073 RTN | | | | | | | |
| 018 ST+0 | | 074 #LBLB | Compute A/G. | | | | | | |
| 019 1 | | 075 RCL | | | | | | | |
| 020 ST+1 | | 076 RCLD | | | | | | | |
| 021 RCL 1 | | 077 RCLC | | | | | | | |
| 022 RTN | | 078 + | | | | | | | |
| 023 #LBLC | Accumulate Σ in R ₀ . | 079 RCLB | $A/G = \frac{\text{Fract}_i}{\sum_{i=2}^5 \text{Fract}_i}$ | | | | | | |
| 024 SPC | | 080 + | | | | | | | |
| 025 RCL i | | 081 RCL A | | | | | | | |
| 026 STOI | I now contains (25 - n). | 082 + | | | | | | | |
| 027 2 | Save in R ₁ . | 083 ÷ | | | | | | | |
| 028 4 | | 084 SPC | | | | | | | |
| 029 STOI | ----- | 085 PRTX | | | | | | | |
| 030 #LBL9 | | 086 RTN | | | | | | | |
| 031 RCL i | | 087 #LBLC | | | | | | | |
| 032 RCL 0 | | 088 INT | Patient ID = Ptnt # .02 | | | | | | |
| 033 ÷ | | 089 . | | | | | | | |
| 034 EE% | $\% = \frac{\text{Fract}_i}{\Sigma} \times 100.$ | 090 0 | | | | | | | |
| 035 2 | | 091 2 | | | | | | | |
| 036 x | | 092 + | | | | | | | |
| 037 PRTX | | 093 STOI | | | | | | | |
| 038 RCL i | | 094 SPC | | | | | | | |
| 039 RCL 1 | Down to R _{25-n} ? | 095 PRTX | | | | | | | |
| 040 X=Y? | | 096 SPC | | | | | | | |
| 041 GTOP | Yes, exit. | 097 RTN | | | | | | | |
| 042 DSZ! | No, decrement and loop again. | 098 #LBLD | | | | | | | |
| 043 GTOP | ----- | 099 F0? | Print toggle. | | | | | | |
| 044 #LBL0 | | 100 GTOP | | | | | | | |
| 045 CLX | Display 0.00 and return. | 101 SF0 | | | | | | | |
| 046 RTN | ----- | 102 1 | | | | | | | |
| 047 #LBLD | Total protein. | 103 RTN | | | | | | | |
| 048 SPC | | 104 #LBL0 | | | | | | | |
| 049 SPC | | 105 CF0 | | | | | | | |
| 050 F0? | | 106 0 | | | | | | | |
| 051 PRTX | | 107 RTN | | | | | | | |
| 052 F0? | | 108 #LBLB | | | | | | | |
| 053 SPC | | 109 2 | | | | | | | |
| 054 STOI | | 110 4 | | | | | | | |
| 055 2 | | 111 STOI | Reprint | | | | | | |
| 056 4 | | 112 SPC | ----- | | | | | | |
| REGISTERS | | | | | | | | | |
| 0 Σ Fract | 1 25 - n | 2 Tot Pr | 3 Ptnt # .02 | 4 | 5 | 6 | 7 | 8 | 9 |
| S0 | S1 | S2 | S3 | S4 | S5 | S6 | S7 | S8 | S9 Fract ₆ |
| A Fract ₅ | B Fract ₄ | C Fract ₃ | D Fract ₂ | E Fract ₁ | F | G | H | I Index | |

| 113 SPC 114 RCL3 115 INT 116 . 117 0 118 2 119 + 120 PRTX 121 SPC 122 #LBL7 123 RCL i 124 PRTX 125 RCL I 126 RCL I 127 X=Y? 128 GT01 129 DSZ I 130 GT07 131 #LBL1 132 2 133 4 134 ST01 135 SPC 136 GSB9 137 SPC 138 SPC 139 RCL2 140 X=0? 141 GT0E 142 PRTX 143 SPC 144 2 145 4 146 ST01 147 GSB8 148 GT0E | Patient ID ----- Loop to print inputs. ----- Print %. If total protein = 0, skip to print A/G. Otherwise print T Pr and grams. Print A/G. | | | | | | | | |
|---|--|------------|------------|-----------|---------|-------|--|------|------------|
| LABELS | | | | | | FLAGS | SET STATUS | | |
| A Start | B Fract | C → % | D T Pr → g | E → A/G | 0 Print | FLAGS | | TRIG | DISP |
| a | b | c Pnt # | d p off? | e Reprint | 1 | 0 | ON OFF | DEG | FIX |
| 0 Used | 1 Used | 2 | 3 | 4 | 2 | 1 | <input type="checkbox"/> <input checked="" type="checkbox"/> | GRAD | SCI |
| 5 | 6 | 7 Prt frac | 8 Prt gms | 9 Prt % | 3 | 2 | <input type="checkbox"/> <input checked="" type="checkbox"/> | RAD | ENG |
| | | | | | | 3 | <input type="checkbox"/> <input checked="" type="checkbox"/> | | n <u>2</u> |

LDH Isoenzymes

| | | | |
|-----------|----------------------------------|------------|--|
| 001 *LBLA | | 057 RTN | |
| 002 2 | Initialize. | 058 *LBL1 | |
| 003 5 | | 059 RCL0 | |
| 004 ST01 | | 060 + | |
| 005 CLX | | 061 EEX | |
| 006 ST00 | | 062 2 | |
| 007 ST01 | | 063 x | |
| 008 ST02 | | 064 PRTX | |
| 009 RTN | | 065 X>Y? | |
| 010 *LBLB | Input LDH values. | 066 R4 | |
| 011 DSZ? | LDH ₁ → R26-1 | 067 SF2 | |
| 012 ST01 | | 068 X>Y? | |
| 013 ST+0 | | 069 SF2 | |
| 014 F0? | Accumulate Σ in R ₀ . | 070 RTN | |
| 015 PRTX | | 071 *LBLC | |
| 016 1 | | 072 INT | |
| 017 ST+1 | | 073 . | |
| 018 RCL1 | Display i. | 074 0 | |
| 019 RTN | | 075 3 | |
| 020 *LBLC | Calculate and print | 076 + | |
| 021 SPC | percentages. | 077 0 | |
| 022 3 | | 078 ST02 | |
| 023 2 | Max LDH ₁ = 33. | 079 PRTX | |
| 024 ENT+ | | 080 SPC | |
| 025 1 | | 081 RTN | |
| 026 0 | Min LDH ₁ = 18. | 082 *LBLD | |
| 027 RCLC | LDH ₁ | 083 F0? | |
| 028 GSB1 | | 084 ST00 | |
| 029 4 | | 085 SF0 | |
| 030 0 | Max LDH ₂ = 40. | 086 1 | |
| 031 ENT+ | | 087 RTN | |
| 032 2 | | 088 *LBL0 | |
| 033 0 | Min LDH ₂ = 28. | 089 CF0 | |
| 034 RCLD | LDH ₂ | 090 0 | |
| 035 GSB1 | | 091 RTN | |
| 036 3 | | 092 *LBL E | |
| 037 0 | Max LDH ₃ = 30. | 093 SPC | |
| 038 ENT+ | | 094 SPC | |
| 039 1 | | 095 RCL2 | |
| 040 0 | Min LDH ₃ = 18. | 096 INT | |
| 041 RCLC | LDH ₃ | 097 0 | |
| 042 GSB1 | | 098 3 | |
| 043 1 | | 099 + | |
| 044 6 | Max LDH ₄ = 16. | 100 PRTX | |
| 045 ENT+ | | 101 SPC | |
| 046 6 | Min LDH ₄ = 6. | 102 RCLC | |
| 047 RCLB | LDH ₄ | 103 PRTX | |
| 048 GSB1 | | 104 RCLD | |
| 049 1 | | 105 PRTX | |
| 050 3 | Max LDH ₅ = 13. | 106 RCLC | |
| 051 ENT+ | | 107 PRTX | |
| 052 2 | Min LDH ₅ = 2. | 108 RCLB | |
| 053 RCLA | LDH ₅ | 109 PRTX | |
| 054 GSE1 | | 110 RCLA | |
| 055 F2? | | 111 FRTX | |
| 056 GTOE | F2 set indicates range error. | | |

Subroutine to find % and test if within normal range.

$$(\%)_i = \frac{LDH_i}{\sum_j LDH_j} \times 100$$

Min > %?

Yes, set flag 2.

% > Max?

Yes, set flag 2.

Patient ID = Ptnt # .03

Print toggle

Reprint

Ptnt # .03

LDH₁

LDH₂

LDH₃

LDH₄

LDH₅

| REGISTERS | | | | | | | | | |
|----------------------|--------------------|--------------------|--------------------|--------------------|-------|----|----|----|----|
| 0 Σ LDH _i | 1 i | 2 Ptnt # .03 | 3 | 4 | 5 | 6 | 7 | 8 | 9 |
| S0 | S1 | S2 | S3 | S4 | S5 | S6 | S7 | S8 | S9 |
| A LDH ₅ | B LDH ₄ | C LDH ₃ | D LDH ₂ | E LDH ₁ | Index | | | | |

Body Surface Area

| | | | | |
|-----------|-----------|---|-----------|--|
| 001 #LBLA | 057 7 | <div>Height (+cm, - in)</div> <div>If height in cm, GTO 1.</div> <div>Convert inches to cm.</div> <div>Store height in cm.</div> <div>Weight (+kg, - lb)</div> <div>If weight in kg, GTO 2.</div> <div>Convert pounds to kg.</div> <div>Store weight in kg.</div> <div>Dubois BSA</div> | 058 1 | <div>Boyd BSA</div> <div>-----</div> <div>Clear for reprint</div> <div>-----</div> <div>Patient ID = Ptnt #.04</div> |
| 002 STOE | 059 8 | | 059 8 | |
| 003 F0? | 060 4 | | 060 4 | |
| 004 SPC | 061 x | | 061 x | |
| 005 F0? | 062 STOA | | 062 STOA | |
| 006 PRTX | 063 F0? | | 063 F0? | |
| 007 X0? | 064 PRTX | | 064 PRTX | |
| 008 GTO1 | 065 F0? | | 065 F0? | |
| 009 CHS | 066 SPC | | 066 SPC | |
| 010 2 | 067 RTM | | 067 RTM | |
| 011 . | 068 #LBLD | | 068 #LBLD | |
| 012 5 | 069 RCLD | | 069 RCLD | |
| 013 4 | 070 . | | 070 . | |
| 014 x | 071 3 | | 071 3 | |
| 015 #LBL1 | 072 Y* | | 072 Y* | |
| 016 STOD | 073 RCLB | | 073 RCLB | |
| 017 RTM | 074 EEX | | 074 EEX | |
| 018 #LBLB | 075 3 | | 075 3 | |
| 019 STOC | 076 x | | 076 x | |
| 020 F0? | 077 ENT† | | 077 ENT† | |
| 021 PRTX | 078 LOE | 078 LOE | | |
| 022 F0? | 079 . | 079 . | | |
| 023 SPC | 080 0 | 080 0 | | |
| 024 X0? | 081 1 | 081 1 | | |
| 025 GTO2 | 082 8 | 082 8 | | |
| 026 CHS | 083 8 | 083 8 | | |
| 027 . | 084 x | 084 x | | |
| 028 4 | 085 . | 085 . | | |
| 029 5 | 086 7 | 086 7 | | |
| 030 3 | 087 2 | 087 2 | | |
| 031 5 | 088 6 | 088 6 | | |
| 032 9 | 089 5 | 089 5 | | |
| 033 2 | 090 - | 090 - | | |
| 034 3 | 091 Y* | 091 Y* | | |
| 035 7 | 092 ÷ | 092 ÷ | | |
| 036 x | 093 3 | 093 3 | | |
| 037 #LBL2 | 094 1 | 094 1 | | |
| 038 STOB | 095 1 | 095 1 | | |
| 039 RTM | 096 8 | 096 8 | | |
| 040 #LBLC | 097 ÷ | 097 ÷ | | |
| 041 RCLD | 098 STOA | 098 STOA | | |
| 042 . | 099 F0? | 099 F0? | | |
| 043 7 | 100 PRTX | 100 PRTX | | |
| 044 2 | 101 F0? | 101 F0? | | |
| 045 5 | 102 SPC | 102 SPC | | |
| 046 Y* | 103 RTM | 103 RTM | | |
| 047 RCLB | 104 #LBL0 | 104 #LBL0 | | |
| 048 . | 105 0 | 105 0 | | |
| 049 4 | 106 STOI | 106 STOI | | |
| 050 2 | 107 RTM | 107 RTM | | |
| 051 5 | 108 #LBLc | 108 #LBLc | | |
| 052 Y* | 109 INT | 109 INT | | |
| 053 x | 110 . | 110 . | | |
| 054 . | 111 0 | 111 0 | | |
| 055 0 | 112 4 | 112 4 | | |
| 056 0 | | | | |

| | | | | | | | | | | | | | | | | | |
|-----------|----------|----|----|---------|----|----|----------|----|----|---------|--|---|----------|--|---|-----------|--|
| REGISTERS | | | | | | | | | | | | | | | | | |
| 0 | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | | | | | | | | |
| S0 | S1 | S2 | S3 | S4 | S5 | S6 | S7 | S8 | S9 | | | | | | | | |
| A | BSA (m²) | | B | Wt (kg) | | C | Wt input | | D | Ht (cm) | | E | Ht input | | F | Ptnt #.04 | |

| | | | | | | | | | |
|-----|-------|--------------|--|--|--|--|--|--|--|
| 113 | + | | | | | | | | |
| 114 | STOI | | | | | | | | |
| 115 | SPC | | | | | | | | |
| 116 | PRTX | | | | | | | | |
| 117 | RTN | | | | | | | | |
| 118 | *LBLd | Print toggle | | | | | | | |
| 119 | F0? | | | | | | | | |
| 120 | GTO0 | | | | | | | | |
| 121 | SF0 | | | | | | | | |
| 122 | 1 | | | | | | | | |
| 123 | RTN | | | | | | | | |
| 124 | *LBL0 | | | | | | | | |
| 125 | CF0 | | | | | | | | |
| 126 | 0 | | | | | | | | |
| 127 | RTN | | | | | | | | |
| 128 | *LBLe | Reprint | | | | | | | |
| 129 | SPC | | | | | | | | |
| 130 | SPC | | | | | | | | |
| 131 | RCL? | | | | | | | | |
| 132 | INT | | | | | | | | |
| 133 | . | | | | | | | | |
| 134 | 0 | | | | | | | | |
| 135 | 4 | | | | | | | | |
| 136 | + | | | | | | | | |
| 137 | PRTX | Patient ID | | | | | | | |
| 138 | SPC | | | | | | | | |
| 139 | RCL | Ht input | | | | | | | |
| 140 | PRTX | | | | | | | | |
| 141 | RCLC | Wt input | | | | | | | |
| 142 | PRTX | | | | | | | | |
| 143 | SPC | | | | | | | | |
| 144 | RCLA | BSA (m²) | | | | | | | |
| 145 | PRTX | | | | | | | | |
| 146 | SPC | | | | | | | | |
| 147 | RTN | | | | | | | | |

| LABELS | | | | | FLAGS | SET STATUS | | | | |
|--------|----------|---|----------|------------|----------|------------|---------|--|---|---|
| A | Ht (+cm) | B | Wt (+kg) | C →Dubois | D →Boyd | E | 0 Print | FLAGS | TRIG | DISP |
| a | Clear | b | | c Pnt # | d P off? | e Reprint | 1 | ON OFF 0 <input checked="" type="checkbox"/> <input type="checkbox"/> | DEG <input checked="" type="checkbox"/> | FIX <input checked="" type="checkbox"/> |
| 0 | P toggle | 1 | Store ht | 2 Store wt | 3 | 4 | 2 | 1 <input type="checkbox"/> <input checked="" type="checkbox"/> | GRAD <input type="checkbox"/> | SCI <input type="checkbox"/> |
| 5 | | 6 | | 7 | 8 | 9 | 3 | 2 <input type="checkbox"/> <input checked="" type="checkbox"/> | RAD <input type="checkbox"/> | ENG <input type="checkbox"/> |
| | | | | | | | | 3 <input type="checkbox"/> <input checked="" type="checkbox"/> | | n 2 |

Urea Clearance

| | | | | |
|-----------|---|-----------|-------------------------|-------------------------------------|
| 001 #LELA | V/t | 057 . | Otherwise have maximum. | |
| 002 ÷ | ----- | 058 3 | | |
| 003 #LBLB | | 059 3 | | |
| 004 STOE | * V | 060 x | | |
| 005 F0? | | 061 ST08 | | |
| 006 SPC | | 062 F0? | | |
| 007 F0? | | 063 PRTX | | |
| 008 PRTX | | 064 RTN | | |
| 009 F2? | If F2 set, must correct | 065 #LBL0 | | Standard |
| 010 GT00 | for BSA. | 066 RCL9 | | |
| 011 GT01 | ----- | 067 1 | | |
| 012 #LBL0 | | 068 . | | |
| 013 1 | | 069 8 | | |
| 014 . | | 070 5 | | |
| 015 7 | | 071 x | | |
| 016 3 | $\dot{V}_{\text{corr}} = \frac{1.73}{\text{BSA}} \dot{V}$ | 072 ST08 | | |
| 017 RCLA | | 073 F0? | | |
| 018 ÷ | | 074 PRTX | | |
| 019 x | ----- | 075 RTN | Clear for reprint. | |
| 020 #LBL1 | Output | 076 #LBL0 | | |
| 021 ST0D | | 077 0 | | |
| 022 F0? | | 078 ST08 | | |
| 023 PRTX | | 079 ST01 | | |
| 024 F0? | | 080 RTN | | |
| 025 SPC | | 081 #LBL6 | | |
| 026 RTN | | 082 SF2 | | Set F2 to allow correction for BSA. |
| 027 #LBLC | | 083 RCLA | | |
| 028 ST0C | U _{urea} | 084 RTN | | |
| 029 F0? | | 085 #LBLC | | |
| 030 PRTX | | 086 INT | Patient ID = Ptnr # .05 | |
| 031 RTN | | 087 . | | |
| 032 #LBLD | | 088 0 | | |
| 033 ST0B | B _{urea} | 089 5 | | |
| 034 F0? | | 090 + | | |
| 035 PRTX | | 091 ST01 | | |
| 036 F0? | | 092 SPC | | |
| 037 SPC | | 093 PRTX | | |
| 038 2 | If $\dot{V}_{\text{corr}} \leq 2$, take $\sqrt{\dot{V}}$ for | 094 RTN | | |
| 039 RCLD | $C_s = \frac{U\sqrt{\dot{V}}}{B}$ | 095 #LBLd | | Print toggle |
| 040 X≠Y? | | 096 F0? | | |
| 041 JX | | 097 GT00 | | |
| 042 RCLC | | 098 SF0 | | |
| 043 RCLB | Otherwise $C_m = \frac{U\dot{V}}{B}$ | 099 1 | | |
| 044 ÷ | | 100 RTN | | |
| 045 x | | 101 #LBL0 | | |
| 046 ST09 | Clearance | 102 CF0 | | |
| 047 F0? | | 103 0 | | |
| 048 PRTX | | 104 RTN | | |
| 049 RTN | | 105 #LBLe | Reprint | |
| 050 #LBLE | % mean normal | 106 SPC | | |
| 051 2 | | 107 SPC | | |
| 052 RCLD | | 108 RCLi | | |
| 053 X≠Y? | If $\dot{V}_{\text{corr}} \leq 2$, GTO 0 for | 109 INT | | |
| 054 GT00 | standard. | 110 . | | |
| 055 RCL9 | | 111 0 | | |
| 056 1 | | 112 5 | | |

| REGISTERS | | | | | | | | | |
|-------------------------|---------------------|---------------------|------------------------------------|----------------------|--------------|----|----|----------|-----|
| 0 | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 % m.n. | 9 C |
| S0 | S1 | S2 | S3 | S4 | S5 | S6 | S7 | S8 | S9 |
| A BSA (m ²) | B B _{urea} | C U _{urea} | D \dot{V}_{corr} (ml/min) | E \dot{V} (ml/min) | I Ptnr # .05 | | | | |

Creatinine Clearance

| | | | | | |
|-----------|---|--|-----------|--|------------------------|
| 001 *LBL# | | | 057 . | | Patient ID = Pnt # .06 |
| 002 ÷ | V/t ----- | | 059 0 | | |
| 003 *LBL# | | | 059 € | | |
| 004 STOE | ṽ | | 060 + | | |
| 005 F0? | | | 061 STOI | | |
| 006 SPC | | | 062 PRTX | | |
| 007 F0? | | | 063 SPC | | |
| 008 PRTX | | | 064 RTN | | |
| 009 F2? | If F2 set, must correct for | | 065 *LBL# | | |
| 010 GT00 | BSA. | | 066 F0? | | Print toggle |
| 011 GT01 | ----- | | 067 GT00 | | |
| 012 *LBL# | | | 068 SPC | | |
| 013 1 | | | 069 1 | | |
| 014 . | | | 070 RTN | | |
| 015 7 | $V_{corr} = \frac{1.73}{BSA} \cdot \dot{V}$ | | 071 *LBL# | | |
| 016 3 | | | 072 CF0 | | |
| 017 RCLA | | | 073 6 | | |
| 018 ÷ | | | 074 RTN | | |
| 019 x | ----- | | 075 *LBL# | | Reprint |
| 020 *LBL# | | | 076 SPC | | |
| 021 STOE | Output | | 077 SPC | | |
| 022 F0? | | | 078 RCL# | | |
| 023 PRTX | | | 079 INT | | |
| 024 F0? | | | 080 . | | |
| 025 SPC | | | 081 0 | | |
| 026 RTN | ----- | | 082 € | | |
| 027 *LBL# | | | 083 + | | |
| 028 STOE | U_{creat} | | 084 PRTX | | Patient ID |
| 029 F0? | | | 085 SPC | | \dot{V} |
| 030 PRTX | ----- | | 086 RCL# | | \dot{V}_{corr} |
| 031 RTN | | | 087 PRTX | | |
| 032 *LBL# | P_{creat} | | 088 RCLD | | |
| 033 STOE | | | 089 PRTX | | |
| 034 F0? | | | 090 SPC | | U_{creat} |
| 035 PRTX | | | 091 RCLD | | P_{creat} |
| 036 RCLC | | | 092 PRTX | | |
| 037 RCLD | $C = \frac{U \cdot \dot{V}}{P}$ | | 093 RCL# | | C |
| 038 x | | | 094 PRTX | | |
| 039 RCL# | | | 095 SPC | | |
| 040 ÷ | | | 096 RCL# | | |
| 041 STOE | | | 097 PRTX | | |
| 042 F0? | | | 098 RTN | | |
| 043 SPC | | | | | |
| 044 F0? | | | | | |
| 045 PRTX | | | | | |
| 046 RTN | | | | | |
| 047 *LBL# | Clear for reprint. | | | | |
| 048 0 | | | | | |
| 049 STOI | | | | | |
| 050 RTN | | | | | |
| 051 *LBL# | | | | | |
| 052 SF2 | Set F2 to allow correction | | | | |
| 053 RCLA | for BSA. | | | | |
| 054 RTN | ----- | | | | |
| 055 *LBL# | | | | | |
| 056 INT | | | | | |

| REGISTERS | | | | | | | | | |
|------------|----------------------|----|----------------------|----|------------------------------|----|--------------|----|-------------|
| 0 | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 C |
| S0 | S1 | S2 | S3 | S4 | S5 | S6 | S7 | S8 | S9 |
| A BSA (m²) | B P _{creat} | | C U _{creat} | | D V _{corr} (ml/min) | | E V (ml/min) | | I Pnt # .06 |

| | | | | | | | | | | | | | | | | | | | |
|---------|------------|----------------------|----------------------|-----------|-----------|---|--|---|--|---|--|------------|--|--|--|--|--|--|--|
| | | | | | | | | | | | | | | | | | | | |
| LABELS | | | | | | | | | | FLAGS | | SET STATUS | | | | | | | |
| A vtt | B V | C U _{creat} | D P _{creat} | E | 0 Print | FLAGS | | TRIG | | DISP | | | | | | | | | |
| a Clear | b Cor BSA? | c Pnt # | d P off? | e Reprint | 1 | ON OFF | | DEG <input checked="" type="checkbox"/> | | FIX <input checked="" type="checkbox"/> | | | | | | | | | |
| 0 Used | 1 Exit V | 2 | 3 | 4 | 2 Cor BSA | 0 <input checked="" type="checkbox"/> 1 <input checked="" type="checkbox"/> 2 <input checked="" type="checkbox"/> 3 <input checked="" type="checkbox"/> | | GRAD <input type="checkbox"/> | | SCI <input type="checkbox"/> | | | | | | | | | |
| 5 | 6 | 7 | 8 | 9 | 3 | | | RAD <input type="checkbox"/> | | ENG <input type="checkbox"/> | | | | | | | | | |
| | | | | | | | | | | | | n <u>2</u> | | | | | | | |

Amniotic Fluid Assay

| | | | |
|--------------------|--|-----------|--------------------------|
| 001 *LBLA | | 057 RCLB | Δ A450 (y) |
| 002 FIX | | 058 XZY | |
| 003 DSP2 | | 059 ÷ | b = y/a ^x |
| 004 STOE | A365 | 060 ST09 | |
| 005 F0? | | 061 F0? | |
| 006 SPC | | 062 PRTX | |
| 007 F0? | | 063 RTN | |
| 008 PRTX | | 064 *LBLC | ----- |
| 009 RTN | | 065 3 | |
| 010 *LBLB | | 066 RCL9 | |
| 011 ST0D | A650 | 067 . | |
| 012 F0? | | 068 7 | |
| 013 PRTX | | 069 XZY? | If b < 0.7, have zone 1. |
| 014 RTN | | 070 GT01 | |
| 015 *LBLC | | 071 R+ | |
| 016 STOC | A450 | 072 XZY? | If b > 3, have zone 3. |
| 017 F0? | | 073 GT03 | |
| 018 PRTX | | 074 2 | Otherwise, have zone 2. |
| 019 RCLC | | 075 GT00 | |
| 020 LN | | 076 *LBL1 | |
| 021 RCLD | | 077 1 | |
| 022 LN | | 078 GT00 | |
| 023 - | | 079 *LBL3 | |
| 024 . | | 080 3 | |
| 025 5 | This changes if different wavelengths of light are used. | 081 *LBL0 | |
| 026 4 | | 082 ST08 | Zone number |
| 027 1 | | 083 FIX | |
| 028 x | | 084 DSP0 | |
| 029 RCLD | | 085 F0? | |
| 030 LN | | 086 PRTX | |
| 031 + | | 087 RTN | |
| 032 e ^x | | 088 *LBLa | ----- |
| 033 - | | 089 CLX | Initialize |
| 034 ST0B | Δ A450 | 090 ST08 | |
| 035 F0? | | 091 ST09 | |
| 036 SPC | | 092 ST0A | |
| 037 F0? | | 093 RTN | |
| 038 PRTX | | 094 *LBLc | |
| 039 F0? | | 095 INT | Patient ID = Ptnt # .07 |
| 040 SPC | | 096 . | |
| 041 RTN | | 097 0 | |
| 042 *LBLD | | 098 7 | |
| 043 ST0A | Week (x) | 099 + | |
| 044 FIX | | 100 ST01 | |
| 045 DSP0 | | 101 PRTX | |
| 046 F0? | | 102 SPC | |
| 047 PRTX | | 103 RTN | |
| 048 DSP2 | | 104 *LBLd | ----- |
| 049 . | | 105 F0? | Print toggle |
| 050 9 | | 106 GT00 | |
| 051 1 | | 107 SF0 | |
| 052 5 | Slope constant a | 108 1 | |
| 053 0 | | 109 RTN | |
| 054 9 | | 110 *LBL0 | |
| 055 XZY | | 111 CF0 | |
| 056 Y* | a ^x | 112 0 | |

| REGISTERS | | | | | | | | | |
|-----------|----------|--------|--------|--------|--------------|----|----|--------|-----|
| 0 | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 Zone | 9 b |
| S0 | S1 | S2 | S3 | S4 | S5 | S6 | S7 | S8 | S9 |
| A Week | B Δ A450 | C A450 | D A550 | E A365 | F Ptnt # .07 | | | | |

| | | | | | |
|-----|--------|------------|--|--|--|
| 113 | RTN | ----- | | | |
| 114 | *LBL e | Reprint | | | |
| 115 | FIX | | | | |
| 116 | DSP2 | | | | |
| 117 | SPC | | | | |
| 118 | SPC | | | | |
| 119 | RCL1 | | | | |
| 120 | INT | | | | |
| 121 | . | | | | |
| 122 | 0 | | | | |
| 123 | 7 | | | | |
| 124 | + | | | | |
| 125 | PRTX | Ptnt # .07 | | | |
| 126 | SPC | | | | |
| 127 | RCL e | A365 | | | |
| 128 | PRTX | | | | |
| 129 | RCLD | A550 | | | |
| 130 | PRTX | | | | |
| 131 | RCLC | A450 | | | |
| 132 | PRTX | | | | |
| 133 | SPC | | | | |
| 134 | RCLB | Δ A450 | | | |
| 135 | PRTX | | | | |
| 136 | SPC | | | | |
| 137 | RCL A | Week | | | |
| 138 | DSP0 | | | | |
| 139 | PRTX | | | | |
| 140 | RCL9 | b | | | |
| 141 | DSP2 | | | | |
| 142 | PRTX | | | | |
| 143 | RCL8 | Zone | | | |
| 144 | DSP0 | | | | |
| 145 | PRTX | | | | |
| 146 | RTN | ----- | | | |

| LABELS | | | | | | FLAGS | SET STATUS | | |
|---------|----------|------------|----------|-----------|---------|-------|--|---|---|
| A A365 | B A550 | C A450 → Δ | D Wk → b | E Zone | 0 Print | | FLAGS | TRIG | DISP |
| a Clear | b | c Ptnt # | d P off? | e Reprint | 1 | | ON OFF | | |
| 0 Used | 1 Zone 1 | 2 | 3 Zone 3 | 4 | 2 | | 0 <input checked="" type="checkbox"/> OFF | DEG <input checked="" type="checkbox"/> | FIX <input checked="" type="checkbox"/> |
| 5 | 6 | 7 | 8 | 9 | 3 | | 1 <input type="checkbox"/> <input checked="" type="checkbox"/> | GRAD <input type="checkbox"/> | SCI <input type="checkbox"/> |
| | | | | | | | 2 <input type="checkbox"/> <input checked="" type="checkbox"/> | RAD <input type="checkbox"/> | ENG <input type="checkbox"/> |
| | | | | | | | 3 <input type="checkbox"/> <input checked="" type="checkbox"/> | | n 2 |

Blood Acid-Base Status

| | | | | | | | | | |
|-----|-------|----------------------------------|-----|--------|--|--|--|--|----------------------------------|
| 001 | *LBLA | BT | 057 | - | | | | | |
| 002 | F0? | | 058 | 10* | | | | | |
| 003 | PRTX | | 059 | . | | | | | |
| 004 | F0? | | 060 | 0 | | | | | |
| 005 | SPC | | 061 | 3 | | | | | |
| 006 | 3 | | 062 | 0 | | | | | |
| 007 | 7 | | 063 | 7 | | | | | |
| 008 | X≠Y | | 064 | x | | | | | |
| 009 | - | | 065 | ST05 | | | | | s(10 ^{pH-pK}) |
| 010 | ST09 | 37-BT | 066 | LSTA | | | | | |
| 011 | SF1 | F1 set for BT. | 067 | + | | | | | |
| 012 | RTN | ----- | 068 | RCLD | | | | | |
| 013 | *LBLB | PCO ₂ | 069 | x | | | | | |
| 014 | ST0E | | 070 | ST0A | | | | | TCO ₂ |
| 015 | F0? | | 071 | F0? | | | | | |
| 016 | PRTX | | 072 | PRTX | | | | | |
| 017 | F1? | To correct for BT, GTO 0. | 073 | CF1 | | | | | Clear flag 1. |
| 018 | GTO0 | | 074 | RTN | | | | | ----- |
| 019 | GTO1 | For 37°, GTO 1. | 075 | *LBL E | | | | | |
| 020 | *LBL0 | ----- | 076 | ST0B | | | | | Hgb |
| 021 | RCL9 | | 077 | F0? | | | | | |
| 022 | . | Correct PCO ₂ to 37°. | 078 | SPC | | | | | |
| 023 | 0 | | 079 | F0? | | | | | |
| 024 | 1 | | 080 | PRTX | | | | | |
| 025 | 9 | | 081 | RCL5 | | | | | |
| 026 | x | | 082 | RCLD | | | | | |
| 027 | 10* | | 083 | x | | | | | |
| 028 | x | ----- | 084 | ST06 | | | | | [HCO ₃ ⁻] |
| 029 | *LBL1 | | 085 | 9 | | | | | |
| 030 | ST0D | PCO ₂ (37°C) | 086 | . | | | | | |
| 031 | RTN | ----- | 087 | 5 | | | | | |
| 032 | *LBLC | pH | 088 | ENT† | | | | | |
| 033 | ST0C | | 089 | 1 | | | | | |
| 034 | F0? | | 090 | . | | | | | |
| 035 | PRTX | | 091 | 6 | | | | | |
| 036 | F1? | To correct for BT, GTO 0. | 092 | 3 | | | | | |
| 037 | GTO0 | | 093 | RCL8 | | | | | |
| 038 | GTO2 | For 37°, GTO 2.----- | 094 | x | | | | | |
| 039 | *LBL0 | | 095 | + | | | | | |
| 040 | RCL9 | | 096 | 7 | | | | | |
| 041 | . | Correct pH to 37° | 097 | . | | | | | |
| 042 | 0 | | 098 | 4 | | | | | |
| 043 | 1 | | 099 | RCL6 | | | | | |
| 044 | 4 | | 100 | - | | | | | |
| 045 | 6 | | 101 | x | | | | | |
| 046 | x | | 102 | - | | | | | |
| 047 | - | ----- | 103 | 2 | | | | | |
| 048 | *LBL2 | | 104 | 4 | | | | | |
| 049 | ST0B | pH (37°C) | 105 | - | | | | | |
| 050 | RTN | ----- | 106 | 1 | | | | | |
| 051 | *LBLD | Compute TCO ₂ . | 107 | RCL8 | | | | | |
| 052 | RCL8 | | 108 | . | | | | | |
| 053 | 6 | | 109 | 0 | | | | | |
| 054 | . | | 110 | 1 | | | | | |
| 055 | 1 | | 111 | 4 | | | | | |
| 056 | 1 | | 112 | 3 | | | | | |

| REGISTERS | | | | | | | | | |
|-----------|------------------|----|----------|----|----------|----|------------------------|----|------------------------|
| 0 | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 |
| S0 | S1 | S2 | S3 | S4 | S5 | S6 | S7 | S8 | S9 |
| A | TCO ₂ | B | pH (37°) | C | pH input | D | PCO ₂ (37°) | E | PCO ₂ input |
| | | | | | | | | | |
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|-----|-------|----------------------------------|--|-----|-------|--|--|----------------------------------|--|
| 113 | X | | | 169 | CHS | | | | |
| 114 | - | | | 170 | PRTX | | | BT | |
| 115 | X | | | 171 | SPC | | | | |
| 116 | STO7 | BE | | 172 | *LBL0 | | | PCO ₂ -input | |
| 117 | F0? | | | 173 | RCL0 | | | | |
| 118 | PRTX | | | 174 | PRTX | | | pH input | |
| 119 | RTN | | | 175 | RCL0 | | | TCO ₂ | |
| 120 | RCL6 | [HCO ₃ ⁻] | | 176 | PRTX | | | | |
| 121 | F0? | | | 177 | RCL4 | | | | |
| 122 | PRTX | | | 178 | PRTX | | | | |
| 123 | RTN | | | 179 | SPC | | | | |
| 124 | *LBL0 | Initialize. | | 180 | RCL8 | | | Hgb | |
| 125 | 0 | | | 181 | PRTX | | | BE | |
| 126 | STO6 | | | 182 | RCL7 | | | | |
| 127 | STO7 | | | 183 | PRTX | | | [HCO ₃ ⁻] | |
| 128 | STO8 | | | 184 | RCL6 | | | | |
| 129 | STO9 | | | 185 | PRTX | | | | |
| 130 | STO1 | | | 186 | RTN | | | | |
| 131 | RTN | | | | | | | | |
| 132 | *LBL0 | | | | | | | | |
| 133 | INT | Patient ID = Pnt # .08. | | | | | | | |
| 134 | . | | | | | | | | |
| 135 | 0 | | | | | | | | |
| 136 | 8 | | | | | | | | |
| 137 | + | | | | | | | | |
| 138 | STO1 | | | | | | | | |
| 139 | PRTX | | | | | | | | |
| 140 | SPC | | | | | | | | |
| 141 | RTN | | | | | | | | |
| 142 | *LBL0 | | | | | | | | |
| 143 | F0? | Print toggle | | | | | | | |
| 144 | GTO0 | | | | | | | | |
| 145 | SF0 | | | | | | | | |
| 146 | 1 | | | | | | | | |
| 147 | RTN | | | | | | | | |
| 148 | *LBL0 | | | | | | | | |
| 149 | CF0 | | | | | | | | |
| 150 | 0 | | | | | | | | |
| 151 | RTN | | | | | | | | |
| 152 | *LBL0 | Reprint | | | | | | | |
| 153 | SPC | | | | | | | | |
| 154 | SPC | | | | | | | | |
| 155 | RCL1 | | | | | | | | |
| 156 | INT | | | | | | | | |
| 157 | . | | | | | | | | |
| 158 | 0 | | | | | | | | |
| 159 | 8 | | | | | | | | |
| 160 | + | | | | | | | | |
| 161 | PRTX | Patient ID | | | | | | | |
| 162 | SPC | | | | | | | | |
| 163 | RCL5 | | | | | | | | |
| 164 | X=0? | If no BT entered, GTO 0. | | | | | | | |
| 165 | GTO0 | | | | | | | | |
| 166 | ? | | | | | | | | |
| 167 | ? | | | | | | | | |
| 168 | - | | | | | | | | |

| LABELS | | | | | FLAGS | SET STATUS | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
|--------|----|---|------------------|---|-------|------------|------------------|---|----------|---|-------|---|----|---|----|---|----|---|----|---|----|---|----|---|----|---|----|---|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----|----|-----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| A | BT | B | PCO ₂ | C | pH | D | TCO ₂ | E | Hgb → BE | 0 | Print | 1 | BT | 2 | BT | 3 | BT | 4 | BT | 5 | BT | 6 | BT | 7 | BT | 8 | BT | 9 | BT | 10 | BT | 11 | BT | 12 | BT | 13 | BT | 14 | BT | 15 | BT | 16 | BT | 17 | BT | 18 | BT | 19 | BT | 20 | BT | 21 | BT | 22 | BT | 23 | BT | 24 | BT | 25 | BT | 26 | BT | 27 | BT | 28 | BT | 29 | BT | 30 | BT | 31 | BT | 32 | BT | 33 | BT | 34 | BT | 35 | BT | 36 | BT | 37 | BT | 38 | BT | 39 | BT | 40 | BT | 41 | BT | 42 | BT | 43 | BT | 44 | BT | 45 | BT | 46 | BT | 47 | BT | 48 | BT | 49 | BT | 50 | BT | 51 | BT | 52 | BT | 53 | BT | 54 | BT | 55 | BT | 56 | BT | 57 | BT | 58 | BT | 59 | BT | 60 | BT | 61 | BT | 62 | BT | 63 | BT | 64 | BT | 65 | BT | 66 | BT | 67 | BT | 68 | BT | 69 | BT | 70 | BT | 71 | BT | 72 | BT | 73 | BT | 74 | BT | 75 | BT | 76 | BT | 77 | BT | 78 | BT | 79 | BT | 80 | BT | 81 | BT | 82 | BT | 83 | BT | 84 | BT | 85 | BT | 86 | BT | 87 | BT | 88 | BT | 89 | BT | 90 | BT | 91 | BT | 92 | BT | 93 | BT | 94 | BT | 95 | BT | 96 | BT | 97 | BT | 98 | BT | 99 | BT | 100 | BT | 101 | BT | 102 | BT | 103 | BT | 104 | BT | 105 | BT | 106 | BT | 107 | BT | 108 | BT | 109 | BT | 110 | BT | 111 | BT | 112 | BT | 113 | BT | 114 | BT | 115 | BT | 116 | BT | 117 | BT | 118 | BT | 119 | BT | 120 | BT | 121 | BT | 122 | BT | 123 | BT | 124 | BT | 125 | BT | 126 | BT | 127 | BT | 128 | BT | 129 | BT | 130 | BT | 131 | BT | 132 | BT | 133 | BT | 134 | BT | 135 | BT | 136 | BT | 137 | BT | 138 | BT | 139 | BT | 140 | BT | 141 | BT | 142 | BT | 143 | BT | 144 | BT | 145 | BT | 146 | BT | 147 | BT | 148 | BT | 149 | BT | 150 | BT | 151 | BT | 152 | BT | 153 | BT | 154 | BT | 155 | BT | 156 | BT | 157 | BT | 158 | BT | 159 | BT | 160 | BT | 161 | BT | 162 | BT | 163 | BT | 164 | BT | 165 | BT | 166 | BT | 167 | BT | 168 | BT | 169 | BT | 170 | BT | 171 | BT | 172 | BT | 173 | BT | 174 | BT | 175 | BT | 176 | BT | 177 | BT | 178 | BT | 179 | BT | 180 | BT | 181 | BT | 182 | BT | 183 | BT | 184 | BT | 185 | BT | 186 | BT | 187 | BT | 188 | BT | 189 | BT | 190 | BT | 191 | BT | 192 | BT | 193 | BT | 194 | BT | 195 | BT | 196 | BT | 197 | BT | 198 | BT | 199 | BT | 200 | BT | 201 | BT | 202 | BT | 203 | BT | 204 | BT | 205 | BT | 206 | BT | 207 | BT | 208 | BT | 209 | BT | 210 | BT | 211 | BT | 212 | BT | 213 | BT | 214 | BT | 215 | BT | 216 | BT | 217 | BT | 218 | BT | 219 | BT | 220 | BT | 221 | BT | 222 | BT | 223 | BT | 224 | BT | 225 | BT | 226 | BT | 227 | BT | 228 | BT | 229 | BT | 230 | BT | 231 | BT | 232 | BT | 233 | BT | 234 | BT | 235 | BT | 236 | BT | 237 | BT | 238 | BT | 239 | BT | 240 | BT | 241 | BT | 242 | BT | 243 | BT | 244 | BT | 245 | BT | 246 | BT | 247 | BT | 248 | BT | 249 | BT | 250 | BT | 251 | BT | 252 | BT | 253 | BT | 254 | BT | 255 | BT | 256 | BT | 257 | BT | 258 | BT | 259 | BT | 260 | BT | 261 | BT | 262 | BT | 263 | BT | 264 | BT | 265 | BT | 266 | BT | 267 | BT | 268 | BT | 269 | BT | 270 | BT | 271 | BT | 272 | BT | 273 | BT | 274 | BT | 275 | BT | 276 | BT | 277 | BT | 278 | BT | 279 | BT | 280 | BT | 281 | BT | 282 | BT | 283 | BT | 284 | BT | 285 | BT | 286 | BT | 287 | BT | 288 | BT | 289 | BT | 290 | BT | 291 | BT | 292 | BT | 293 | BT | 294 | BT | 295 | BT | 296 | BT | 297 | BT | 298 | BT | 299 | BT | 300 | BT | 301 | BT | 302 | BT | 303 | BT | 304 | BT | 305 | BT | 306 | BT | 307 | BT | 308 | BT | 309 | BT | 310 | BT | 311 | BT | 312 | BT | 313 | BT | 314 | BT | 315 | BT | 316 | BT | 317 | BT | 318 | BT | 319 | BT | 320 | BT | 321 | BT | 322 | BT | 323 | BT | 324 | BT | 325 | BT | 326 | BT | 327 | BT | 328 | BT | 329 | BT | 330 | BT | 331 | BT | 332 | BT | 333 | BT | 334 | BT | 335 | BT | 336 | BT | 337 | BT | 338 | BT | 339 | BT | 340 | BT | 341 | BT | 342 | BT | 343 | BT | 344 | BT | 345 | BT | 346 | BT | 347 | BT | 348 | BT | 349 | BT | 350 | BT | 351 | BT | 352 | BT | 353 | BT | 354 | BT | 355 | BT | 356 | BT | 357 | BT | 358 | BT | 359 | BT | 360 | BT | 361 | BT | 362 | BT | 363 | BT | 364 | BT | 365 | BT | 366 | BT | 367 | BT | 368 | BT | 369 | BT | 370 | BT | 371 | BT | 372 | BT | 373 | BT | 374 | BT | 375 | BT | 376 | BT | 377 | BT | 378 | BT | 379 | BT | 380 | BT | 381 | BT | 382 | BT | 383 | BT | 384 | BT | 385 | BT | 386 | BT | 387 | BT | 388 | BT | 389 | BT | 390 | BT | 391 | BT | 392 | BT | 393 | BT | 394 | BT | 395 | BT | 396 | BT | 397 | BT | 398 | BT | 399 | BT | 400 | BT | 401 | BT | 402 | BT | 403 | BT | 404 | BT | 405 | BT | 406 | BT | 407 | BT | 408 | BT | 409 | BT | 410 | BT | 411 | BT | 412 | BT | 413 | BT | 414 | BT | 415 | BT | 416 | BT | 417 | BT | 418 | BT | 419 | BT | 420 | BT | 421 | BT | 422 | BT | 423 | BT | 424 | BT | 425 | BT | 426 | BT | 427 | BT | 428 | BT | 429 | BT | 430 | BT | 431 | BT | 432 | BT | 433 | BT | 434 | BT | 435 | BT | 436 | BT | 437 | BT | 438 | BT | 439 | BT | 440 | BT | 441 | BT | 442 | BT | 443 | BT | 444 | BT | 445 | BT | 446 | BT | 447 | BT | 448 | BT | 449 | BT | 450 | BT | 451 | BT | 452 | BT | 453 | BT | 454 | BT | 455 | BT | 456 | BT | 457 | BT | 458 | BT | 459 | BT | 460 | BT | 461 | BT | 462 | BT | 463 | BT | 464 | BT | 465 | BT | 466 | BT | 467 | BT | 468 | BT | 469 | BT | 470 | BT | 471 | BT | 472 | BT | 473 | BT | 474 | BT | 475 | BT | 476 | BT | 477 | BT | 478 | BT | 479 | BT | 480 | BT | 481 | BT | 482 | BT | 483 | BT | 484 | BT | 485 | BT | 486 | BT | 487 | BT | 488 | BT | 489 | BT | 490 | BT | 491 | BT | 492 | BT | 493 | BT | 494 | BT | 495 | BT | 496 | BT | 497 | BT | 498 | BT | 499 | BT | 500 | BT | 501 | BT | 502 | BT | 503 | BT | 504 | BT | 505 | BT | 506 | BT | 507 | BT | 508 | BT | 509 | BT | 510 | BT | 511 | BT | 512 | BT | 513 | BT | 514 | BT | 515 | BT | 516 | BT | 517 | BT | 518 | BT | 519 | BT | 520 | BT | 521 | BT | 522 | BT | 523 | BT | 524 | BT | 525 | BT | 526 | BT | 527 | BT | 528 | BT | 529 | BT | 530 | BT | 531 | BT | 532 | BT | 533 | BT | 534 | BT | 535 | BT | 536 | BT | 537 | BT | 538 | BT | 539 | BT | 540 | BT | 541 | BT | 542 | BT | 543 | BT | 544 | BT | 545 | BT | 546 | BT | 547 | BT | 548 | BT | 549 | BT | 550 | BT | 551 | BT | 552 | BT | 553 | BT | 554 | BT | 555 | BT | 556 | BT | 557 | BT | 558 | BT | 559 | BT | 560 | BT | 561 | BT | 562 | BT | 563 | BT |

Oxygen Saturation and Content

| | | | | | | | | | | |
|---------------------------|--------------------------------------|-----------|--------------------------------|--------------------|----|-------------------------|-------|-------------------------|---------|-------------|
| 001 *LBLA | BT | 057 GT08 | ----- | | | | | | | |
| 002 F0? | | 058 *LBL1 | If input < 0, make positive | | | | | | | |
| 003 PRTX | | 059 CHS | ----- | | | | | | | |
| 004 3 | | 060 *LBL0 | | | | | | | | |
| 005 7 | | 061 ST0C | VPO ₂ | | | | | | | |
| 006 XZY | | 062 ENT+ | | | | | | | | |
| 007 - | | 063 ENT+ | | | | | | | | |
| 008 ST09 | 37-BT | 064 ENT+ | | | | | | | | |
| 009 RCLD | Rcl PCO ₂ (if input). | 065 1 | | | | | | | | |
| 010 RTW | ----- | 066 5 | | | | | | | | |
| 011 *LBLB | PCO ₂ | 067 - | | | | | | | | |
| 012 F0? | | 068 x | | | | | | | | |
| 013 PRTX | | 069 2 | | | | | | | | |
| 014 ST0D | | 070 0 | | | | | | | | |
| 015 RCLB | Rcl pH (if input). | 071 4 | | | | | | | | |
| 016 RTW | ----- | 072 5 | | | | | | | | |
| 017 *LBLC | pH | 073 + | | | | | | | | |
| 018 F0? | | 074 x | | | | | | | | |
| 019 PRTX | | 075 2 | | | | | | | | |
| 020 ST0B | | 076 EE* | Compute oxygen | | | | | | | |
| 021 RTW | | 077 3 | saturation. | | | | | | | |
| 022 *LBLD | ----- | 078 + | | | | | | | | |
| 023 F0? | | 079 x | | | | | | | | |
| 024 PRTX | PO ₂ input | 080 ST07 | | | | | | | | |
| 025 ST0E | If input < 0, consider as | 081 CLX | | | | | | | | |
| 026 X0? | VPO ₂ . | 082 1 | | | | | | | | |
| 027 GT01 | Otherwise compute VPO ₂ . | 083 5 | | | | | | | | |
| 028 RCL9 | | 084 - | | | | | | | | |
| 029 . | | 085 x | | | | | | | | |
| 030 0 | | 086 2 | | | | | | | | |
| 031 2 | | 087 4 | | | | | | | | |
| 032 4 | | 088 0 | | | | | | | | |
| 033 x | | 089 0 | | | | | | | | |
| 034 RCLB | | 090 + | | | | | | | | |
| 035 7 | | 091 x | | | | | | | | |
| 036 . | | 092 3 | | | | | | | | |
| 037 4 | | 093 1 | | | | | | | | |
| 038 - | | 094 1 | | | | | | | | |
| 039 . | | 095 0 | | | | | | | | |
| 040 4 | | 096 0 | | | | | | | | |
| 041 0 | | 097 - | | | | | | | | |
| 042 x | | 098 x | | | | | | | | |
| 043 + | | 099 2 | | | | | | | | |
| 044 4 | | 100 4 | | | | | | | | |
| 045 0 | | 101 EE* | | | | | | | | |
| 046 RCLD | | 102 5 | | | | | | | | |
| 047 ÷ | | 103 + | | | | | | | | |
| 048 LOG | | 104 EE* | | | | | | | | |
| 049 . | | 105 2 | | | | | | | | |
| 050 0 | | 106 ÷ | | | | | | | | |
| 051 6 | | 107 ST÷7 | | | | | | | | |
| 052 x | | 108 RCL7 | O ₂ saturation (%). | | | | | | | |
| 053 + | | 109 SF2 | F2 set to indicate | | | | | | | |
| 054 10* | | 110 F0? | saturation computed. | | | | | | | |
| 055 RCLE | | 111 SPC | | | | | | | | |
| 056 * | | 112 F0? | | | | | | | | |
| REGISTERS | | | | | | | | | | |
| 0 | 1 | 2 | 3 | 4 | 5 | 6 | 7 Sat | 8 Hgb | 9 37-BT | |
| S0 | S1 | S2 | S3 | S4 | S5 | S6 | S7 | S8 | S9 | |
| A. O ₂ content | | B pH(37) | | C VPO ₂ | | D PCO ₂ (37) | | E PO ₂ input | | I Pnt # .09 |

| | | | | | | | |
|-----|-------|--------------------------|--|-----|-------|--|-----------------------|
| 113 | PRTX | | | 169 | PRTX | | |
| 114 | F0? | | | 170 | SPC | | |
| 115 | SPC | | | 171 | STO1 | | |
| 116 | RTN | | | 172 | RTN | | |
| 117 | RCL8 | Rcl Hgb (if input). | | 173 | *LBLd | | Print toggle |
| 118 | R/S | ----- | | 174 | F0? | | |
| 119 | *LBLc | | | 175 | GT00 | | |
| 120 | F2? | If Sat. computed, do not | | 176 | SF0 | | |
| 121 | GT00 | input it. | | 177 | 1 | | |
| 122 | XZY | | | 178 | RTN | | |
| 123 | STO7 | Otherwise store Sat. | | 179 | *LBL0 | | |
| 124 | F0? | | | 180 | CF0 | | |
| 125 | PRTX | | | 181 | 0 | | |
| 126 | F0? | | | 182 | RTN | | |
| 127 | SPC | | | 183 | *LBLc | | Reprint |
| 128 | XZY | ----- | | 184 | SPC | | |
| 129 | *LBL0 | | | 185 | SPC | | |
| 130 | STO8 | Store Hgb. | | 186 | RCL1 | | |
| 131 | F0? | | | 187 | INT | | |
| 132 | PRTX | | | 188 | . | | |
| 133 | RCL7 | | | 189 | 0 | | |
| 134 | x | | | 190 | 9 | | |
| 135 | 1 | | | 191 | + | | |
| 136 | 3 | | | 192 | PRTX | | Patient ID |
| 137 | 4 | | | 193 | SPC | | |
| 138 | x | Compute oxygen content. | | 194 | 3 | | |
| 139 | RCL0 | | | 195 | 7 | | |
| 140 | 3 | | | 196 | RCL9 | | |
| 141 | 1 | | | 197 | - | | |
| 142 | x | | | 198 | PRTX | | BT |
| 143 | + | | | 199 | RCLD | | |
| 144 | EEY | | | 200 | PRTX | | PCO ₂ |
| 145 | 4 | | | 201 | RCLB | | |
| 146 | ÷ | | | 202 | PRTX | | pH |
| 147 | STO0 | O ₂ content | | 203 | RCLc | | |
| 148 | F0? | | | 204 | PRTX | | PO ₂ input |
| 149 | PRTX | | | 205 | SPC | | |
| 150 | RTN | | | 206 | RCL7 | | |
| 151 | *LBLc | ----- | | 207 | PRTX | | Saturation |
| 152 | 0 | Initialize | | 208 | SPC | | |
| 153 | STO0 | | | 209 | RCL8 | | Hemoglobin |
| 154 | STO0 | | | 210 | PRTX | | |
| 155 | STO1 | | | 211 | RCL0 | | Content |
| 156 | RTN | | | 212 | PRTX | | |
| 157 | *LBLb | ----- | | 213 | RTN | | |
| 158 | 3 | | | | | | |
| 159 | 7 | Rcl BT | | | | | |
| 160 | RCL9 | | | | | | |
| 161 | - | | | | | | |
| 162 | RTN | BT = 37 - (37-BT) | | | | | |
| 163 | *LBLc | ----- | | | | | |
| 164 | INT | | | | | | |
| 165 | . | | | | | | |
| 166 | 0 | Patient ID = Pnt # .09 | | | | | |
| 167 | 9 | | | | | | |
| 168 | + | | | | | | |

| LABELS | | | | | FLAGS | SET STATUS | | |
|--------|------------------|-------|-----------------------|-------------------------|--------------|--|---|---|
| A | B | C | D | E | 0 | 1 | 2 | 3 |
| BT | PCO ₂ | pH | PO ₂ → Sat | St Hgb → O ₂ | Print | ON OFF | TRIG | DISP |
| Clear | → Rcl BT | Pnt # | P off? | Reprint | | 0 <input checked="" type="checkbox"/> <input type="checkbox"/> | DEG <input checked="" type="checkbox"/> | FIX <input checked="" type="checkbox"/> |
| Used | VPO ₂ | | | | Sat computed | 1 <input type="checkbox"/> <input checked="" type="checkbox"/> | GRAD <input type="checkbox"/> | SCI <input type="checkbox"/> |
| | | | | | | 2 <input type="checkbox"/> <input checked="" type="checkbox"/> | RAD <input type="checkbox"/> | ENG <input type="checkbox"/> |
| | | | | | | 3 <input type="checkbox"/> <input checked="" type="checkbox"/> | | n <u>2</u> |

Red Cell Indices

| | | | | | | | | | |
|---|---|--|--|---------|--------------|----|----|----|--------|
| 001 *LBLA 002 STOE 003 F0? 004 PRTX 005 RTN 006 *LBLB 007 STOD 008 F0? 009 PRTX 010 RTN 011 *LBLC 012 STOC 013 F0? 014 PRTX 015 F0? 016 SPC 017 RCLD 018 1 019 0 020 x 021 RCLE 022 + 023 STOB 024 F0? 025 PRTX 026 RTN 027 *LBLD 028 RCLC 029 1 030 0 031 x 032 RCLE 033 + 034 STOA 035 F0? 036 PRTX 037 RTN 038 *LBLE 039 RCLC 040 EEX 041 2 042 x 043 RCLD 044 + 045 STOG 046 F0? 047 PRTX 048 RTN 049 *LBLA 050 0 051 STOI 052 RTN 053 *LBLC 054 INT 055 . 056 1 | Count. ----- Hematocrit (%). ----- Hemoglobin. ----- Compute MCV. ----- MCV. ----- Compute MCH. ----- MCH. ----- Compute MCHC. ----- MCHC. ----- Clear. ----- Patient ID = Ptnt # .10 | 057 + 058 STOI 059 PRTX 060 SPC 061 RTN 062 *LBLD 063 F0? 064 STOB 065 SF0 066 1 067 RTN 068 *LBLB 069 CF0 070 0 071 RTN 072 *LBLC 073 SPC 074 SPC 075 RCLC 076 INT 077 . 078 1 079 + 080 PRTX 081 SPC 082 RCLE 083 PRTX 084 RCLD 085 PRTX 086 RCLC 087 PRTX 088 SPC 089 RCLB 090 PRTX 091 RCLC 092 PRTX 093 RCLC 094 PRTX 095 RTN | ----- Print toggle. ----- Reprint. ----- Patient ID. Count. Hct (%) Hgb MCV MCH MCHC ----- | | | | | | |
| REGISTERS | | | | | | | | | |
| 0 | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 MCHC |
| S0 | S1 | S2 | S3 | S4 | S5 | S6 | S7 | S8 | S9 |
| A MCH | B MCV | C Hgb | D Hct (%) | E Count | I Ptnt # .10 | | | | |

| | | | | | | | | | | | | | | | | | | | | | | | |
|---------|--|-----------|--|-----------|--|----------|--|-----------|--|---------|--|---|--|--|--|---|--|--|--|---|--|--|--|
| | | | | | | | | | | | | | | | | | | | | | | | |
| LABELS | | | | | | | | | | | | FLAGS | | | | SET STATUS | | | | | | | |
| A Count | | B Hct (%) | | C Hgb→MCV | | D →MCH | | E →MCHC | | 0 Print | | FLAGS | | | | TRIG | | | | DISP | | | |
| a Clear | | b | | c Ptnt # | | d P off? | | e Reprint | | 1 | | 0 <input checked="" type="checkbox"/> ON <input type="checkbox"/> OFF | | | | DEG <input checked="" type="checkbox"/> | | | | FIX <input checked="" type="checkbox"/> | | | |
| 0 Used | | 1 | | 2 | | 3 | | 4 | | 2 | | 1 <input type="checkbox"/> <input checked="" type="checkbox"/> | | | | GRAD <input type="checkbox"/> | | | | SCI <input type="checkbox"/> | | | |
| 5 | | 6 | | 7 | | 8 | | 9 | | 3 | | 2 <input type="checkbox"/> <input checked="" type="checkbox"/> | | | | RAD <input type="checkbox"/> | | | | ENG <input type="checkbox"/> | | | |
| | | | | | | | | | | | | 3 <input type="checkbox"/> <input checked="" type="checkbox"/> | | | | n <u>2</u> | | | | | | | |

Total Blood Volume

| | | | |
|------------|------------------------------------|-----------|--------------------|
| 001 #LBLA | | 057 #LBLd | Print toggle |
| 002 STOE | Background counts. | 058 F0? | |
| 003 F0? | | 059 GT00 | |
| 004 PRTX | | 060 SF0 | |
| 005 RTN | ----- | 061 1 | |
| 006 #LBLB | Volume injected. | 062 RTN | |
| 007 STOD | | 063 #LBL0 | |
| 008 F0? | | 064 CF0 | |
| 009 PRTX | ----- | 065 0 | |
| 010 RTN | | 066 RTN | ----- |
| 011 #LBLC | Standard dilution. | 067 #LBLc | Reprint |
| 012 STOC | | 068 SPC | |
| 013 F0? | | 069 SPC | |
| 014 PRTX | ----- | 070 RCL! | |
| 015 RTN | | 071 INT | |
| 016 #LBLD | Standard CPM. | 072 . | |
| 017 STOB | | 073 1 | |
| 018 F0? | | 074 1 | |
| 019 PRTX | ----- | 075 + | |
| 020 RTN | | 076 PRTX | Patient ID |
| 021 #LBL E | Whole blood CPM. | 077 SPC | Bck |
| 022 STOA | | 078 RCL | Vol. injected |
| 023 F0? | | 079 PRTX | Std. dilution |
| 024 PRTX | | 080 RCLD | Std. CPM |
| 025 F0? | Net Std. CPM = Std. CPM- Bck. | 081 PRTX | Blood CPM |
| 026 SPC | | 082 RCLC | |
| 027 RCLB | | 083 PRTX | |
| 028 RCLC | | 084 RCLB | |
| 029 - | | 085 PRTX | |
| 030 XZY | Net blood CPM = Blood CPM- Bck. | 086 RCL A | |
| 031 RCL | | 087 PRTX | |
| 032 - | | 088 SPC | |
| 033 ÷ | | 089 RCL9 | Total blood volume |
| 034 RCLC | | 090 PRTX | ----- |
| 035 x | | 091 RTN | |
| 036 RCLD | Total blood volume. | | |
| 037 x | | | |
| 038 ST09 | | | |
| 039 F0? | | | |
| 040 PRTX | ----- | | |
| 041 RTN | Initialize. | | |
| 042 #LBL a | | | |
| 043 0 | | | |
| 044 STOE | | | |
| 045 STOI | ----- | | |
| 046 RTN | Patient ID = Ptnt # .11 | | |
| 047 #LBLc | | | |
| 048 INT | | | |
| 049 . | | | |
| 050 1 | | | |
| 051 1 | | | |
| 052 + | | | |
| 053 STOI | | | |
| 054 PRTX | | | |
| 055 SPC | | | |
| 056 RTN | | | |

| REGISTERS | | | | | | | | | |
|-------------|------------|-----------------|-----------------|-------|--------------|----|----|----|-------|
| 0 | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 TBV |
| S0 | S1 | S2 | S3 | S4 | S5 | S6 | S7 | S8 | S9 |
| A Blood CPM | B Std. CPM | C Std. dilution | D Vol. injected | E Bck | I Ptnt # .11 | | | | |

Schilling Test

| | | | | | | | | | | |
|---|--|--|---|------------|----|-----------------|----|-------|-----|--------------|
| 001 #LBLA 002 STOE 003 F0? 004 PRTX 005 RTN 006 #LBLB 007 STOD 008 F0? 009 PRTX 010 RTN 011 #LBLC 012 STOC 013 F0? 014 PRTX 015 RTN 016 #LBLD 017 STOB 018 F0? 019 PRTX 020 RTN 021 #LBLE 022 STOA 023 F0? 024 PRTX 025 1 026 XZY 027 RCLE 028 - 029 1 030 RCLB 031 XZY? 032 R4 033 x 034 x 035 RCLC 036 RCLE 037 - 038 ÷ 039 RCLD 040 ÷ 041 EEX 042 2 043 x 044 STOG 045 F0? 046 SPC 047 F0? 048 PRTX 049 RTN 050 #LBL0 051 0 052 STOE 053 STOI 054 RTN 055 #LBLC 056 INT | Background counts. ----- Standard dilution. ----- Standard counts. ----- Urine volume. (V) ----- Urine counts. (U) 1 U U 1 Bck U 1 Net 1 1 Net 1 V 1 Net 1 Is V ≤ 1? Yes, eliminate V. No, V > 1, multiply by V. Net std. counts. Convert to %. % of dose excreted. ----- Initialize. ----- | 057 . 058 1 059 2 060 + 061 STOI 062 PRTX 063 SPC 064 RTN 065 #LBL4 066 F0? 067 STOB 068 SFO 069 1 070 RTN 071 #LBL0 072 CFE 073 0 074 RTN 075 #LBLE 076 SPC 077 SPC 078 RCLI 079 INT 080 . 081 1 082 2 083 + 084 PRTX 085 SPC 086 RCLE 087 PRTX 088 RCLD 089 PRTX 090 RCLC 091 PRTX 092 RCLB 093 PRTX 094 RCLA 095 PRTX 096 SPC 097 RCL0 098 PRTX 099 RTN | Patient ID = Ptnt # .12 ----- Print toggle ----- Reprint ----- Patient ID Bck Std. dilution Std. CPM Urine vol. Urine CPM % excreted ----- | | | | | | | |
| REGISTERS | | | | | | | | | | |
| 0 | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 % | |
| S0 | S1 | S2 | S3 | S4 | S5 | S6 | S7 | S8 | S9 | |
| A Urine CPM | | B Urine Vol. | | C Std. CPM | | D Std. dilution | | E Bck | | I Ptnt # .12 |

Thyroid Uptake

| | | | |
|-----------|----------------------------|-----------|--|
| 001 #LBLA | Standard Counts. | 057 RCLC | |
| 002 STOE | | 058 RCLB | |
| 003 F0? | | 059 - | |
| 004 PRTY | | 060 + | |
| 005 RTN | ----- | 061 LSTX | |
| 006 #LBLB | | 062 ÷ | |
| 007 STOD | Standard Background. | 063 STX9 | |
| 008 F0? | | 064 RCL9 | |
| 009 PRTX | | 065 F0? | |
| 010 RCLC | | 066 PRTX | |
| 011 XZY | | 067 RTN | |
| 012 - | Std. Cts. - Std. Bck. | 068 #LBLB | |
| 013 RTN | ----- | 069 F0? | |
| 014 #LBLC | | 070 SPC | |
| 015 STOC | Patient Counts. | 071 XZY | |
| 016 F0? | | 072 F0? | |
| 017 PRTX | | 073 PRTX | |
| 018 RTN | ----- | 074 XZY | |
| 019 #LBLD | | 075 F0? | |
| 020 STOB | Patient Background. | 076 PRTX | |
| 021 F0? | | 077 ÷ | |
| 022 PRTX | | 078 STX9 | |
| 023 RCLC | | 079 RCL9 | |
| 024 XZY | | 080 F0? | |
| 025 - | Ptnt. Cts. - Ptnt. Bck. | 081 PRTX | |
| 026 RTN | ----- | 082 RTN | |
| 027 #LBLC | Compute uptake. | 083 #LBLC | |
| 028 RCLC | | 084 INT | |
| 029 RCLB | | 085 . | |
| 030 - | | 086 1 | |
| 031 RCLC | | 087 3 | |
| 032 RCLD | | 088 + | |
| 033 - | | 089 STOI | |
| 034 ÷ | | 090 PRTX | |
| 035 EEX | | 091 SPC | |
| 036 2 | | 092 RTN | |
| 037 x | | 093 #LBLD | |
| 038 STOD | % Uptake. | 094 F0? | |
| 039 F0? | | 095 STOB | |
| 040 SPC | | 096 SFO | |
| 041 F0? | | 097 1 | |
| 042 PRTX | | 098 RTN | |
| 043 RTN | | 099 #LBLB | |
| 044 #LBLA | Correction for prior | 100 CFB | |
| 045 F0? | radioactivity. | 101 0 | |
| 046 SPC | | 102 RTN | |
| 047 XZY | | 103 #LBLC | |
| 048 F0? | | 104 SPC | |
| 049 PRTX | Patient predose counts. | 105 SPC | |
| 050 XZY | | 106 RCLC | |
| 051 F0? | | 107 INT | |
| 052 PRTX | Background predose counts. | 108 . | |
| 053 - | | 109 1 | |
| 054 RCLC | Decay factor. | 110 3 | |
| 055 x | | 111 + | |
| 056 CHS | | 112 PRTX | |

| REGISTERS | | | | | | | | | |
|----------------|----|-------------|----|--------------|----|------------|----|--------------|------------|
| 0 | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 % Uptake |
| S0 | S1 | S2 | S3 | S4 | S5 | S6 | S7 | S8 | S9 |
| A Decay factor | | B Ptnt. Bck | | C Ptnt. Cts. | | D Std. Bck | | E Std. Cts. | |
| | | | | | | | | F Ptnt # .13 | |

[illegible]

Radioactive Decay Corrections

| | | | | | | | | | | | |
|------------------------------------|--|------------------------|----------------------------------|-----|----|-------------|----|------------------|----|---------------------|--|
| 001 #LBLA | Initial activity (A ₀). | 057 STOD | Store t (hrs). | | | | | | | | |
| 002 F3? | | 058 RCLB | | | | | | | | | |
| 003 STOB | | 059 ÷ | | | | | | | | | |
| 004 RCLC | Calculate: A ₀ = A/f | 060 . | Store decay factor | | | | | | | | |
| 005 RCLA | | 061 5 | | | | | | | | | |
| 006 ÷ | | 062 XZY | $f = \frac{1}{2}^{t/\tau_{1/2}}$ | | | | | | | | |
| 007 STOE | | 063 Y* | | | | | | | | | |
| 008 RTN | ----- | 064 STOW | f = $\frac{1}{2}$ | | | | | | | | |
| 009 #LBL0 | Store input A ₀ . | 065 R† | Display t as input. | | | | | | | | |
| 010 STOE | ----- | 066 RTN | ----- | | | | | | | | |
| 011 RTN | Time in days, hours. | 067 #LBLC | Present activity (A). | | | | | | | | |
| 012 #LBLB | | 068 F3? | | | | | | | | | |
| 013 F3? | | 069 STOB | | | | | | | | | |
| 014 STOB | | 070 RCLC | Calculate: | | | | | | | | |
| 015 RCLC | | 071 RCLA | A = A ₀ f | | | | | | | | |
| 016 RCLC | | 072 x | ----- | | | | | | | | |
| 017 ÷ | Calculate: | 073 STOC | | | | | | | | | |
| 018 STOW | | 074 RTN | | | | | | | | | |
| 019 LN | $t = \frac{\tau_{1/2} \ln f}{\ln 1/2}$ | 075 #LBL0 | Store input A. | | | | | | | | |
| 020 . | | 076 STOC | ----- | | | | | | | | |
| 021 5 | | 077 RTN | Isotopes 1–9. | | | | | | | | |
| 022 LN | | 078 #LBLD | | | | | | | | | |
| 023 ÷ | | 079 STOI | | | | | | | | | |
| 024 RCLB | | 080 GSB† | Store $\tau_{1/2}$. | | | | | | | | |
| 025 x | Store t (hours). | 081 STOB | ----- | | | | | | | | |
| 026 STOD | | 082 RTN | | | | | | | | | |
| 027 2 | | 083 #LBLC | | | | | | | | | |
| 028 4 | | 084 6 | | | | | | | | | |
| 029 ÷ | | 085 6 | | | | | | | | | |
| 030 INT | Convert t in hrs. to dd.hh | 086 7 | ⁵¹ Cr | | | | | | | | |
| 031 ENT† | for display. | 087 . | | | | | | | | | |
| 032 ENT† | | 088 2 | | | | | | | | | |
| 033 2 | | 089 STOB | | | | | | | | | |
| 034 4 | | 090 RTN | | | | | | | | | |
| 035 x | | 091 #LBLB | ----- | | | | | | | | |
| 036 RCLD | | 092 6 | | | | | | | | | |
| 037 XZY | | 093 4 | | | | | | | | | |
| 038 - | | 094 6 | ⁵⁷ Co | | | | | | | | |
| 039 EEX | | 095 0 | | | | | | | | | |
| 040 2 | | 096 STOB | | | | | | | | | |
| 041 ÷ | | 097 RTN | | | | | | | | | |
| 042 + | | 098 #LBLC | | | | | | | | | |
| 043 RTN | ----- | 099 6 | ^{99m} Tc | | | | | | | | |
| 044 #LBL0 | Time input. | 100 STOB | ----- | | | | | | | | |
| 045 ENT† | | 101 RTN | | | | | | | | | |
| 046 ENT† | | 102 #LBLD | | | | | | | | | |
| 047 INT | | 103 1 | | | | | | | | | |
| 048 2 | | 104 4 | | | | | | | | | |
| 049 4 | Convert from dd.hh | 105 4 | | | | | | | | | |
| 050 x | format to hours. | 106 0 | ¹²⁵ I | | | | | | | | |
| 051 XZY | | 107 STOB | | | | | | | | | |
| 052 FRC | | 108 RTN | | | | | | | | | |
| 053 EEX | | 109 #LBLC | ----- | | | | | | | | |
| 054 2 | | 110 1 | | | | | | | | | |
| 055 x | | 111 9 | | | | | | | | | |
| 056 + | | 112 3 | ¹³¹ I | | | | | | | | |
| REGISTERS | | | | | | | | | | | |
| 0 | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | | |
| S0 | S1 | S2 | S3 | S4 | S5 | S6 | S7 | S8 | S9 | | |
| A Decay factor (A/A ₀) | | B $\tau_{1/2}$ (hours) | | C A | | D t (hours) | | E A ₀ | | I Isotope no. (1–9) | |

| | | | |
|-----------|-------|-----------|--------|
| 113 . | | 169 . | |
| 114 2 | | 170 7 | |
| 115 STOP | | 171 3 | 113mIn |
| 116 RTN | ----- | 172 PTM | |
| 117 #LBL1 | | 173 #LBL8 | |
| 118 2 | | 174 1 | |
| 119 6 | 137Cs | 175 2 | 133Xe |
| 120 2 | | 176 6 | |
| 121 9 | | 177 . | |
| 122 8 | | 178 5 | |
| 123 0 | | 179 RTN | |
| 124 STOP | ----- | 180 #LBL9 | |
| 125 RTN | | 181 6 | 197Hg |
| 126 #LBL1 | | 182 5 | |
| 127 1 | | 183 RTN | |
| 128 0 | | | |
| 129 7 | 3H | | |
| 130 4 | | | |
| 131 7 | | | |
| 132 0 | | | |
| 133 RTN | ----- | | |
| 134 #LBL2 | | | |
| 135 5 | | | |
| 136 0 | 14C | | |
| 137 5 | | | |
| 138 8 | | | |
| 139 EEY | | | |
| 140 4 | | | |
| 141 RTN | ----- | | |
| 142 #LBL3 | | | |
| 143 1 | | | |
| 144 . | 18F | | |
| 145 8 | | | |
| 146 7 | | | |
| 147 RTN | ----- | | |
| 148 #LBL4 | | | |
| 149 3 | 32P | | |
| 150 4 | | | |
| 151 3 | | | |
| 152 . | | | |
| 153 2 | | | |
| 154 RTN | ----- | | |
| 155 #LBL5 | | | |
| 156 2 | | | |
| 157 8 | 75Se | | |
| 158 8 | | | |
| 159 0 | | | |
| 160 RTN | ----- | | |
| 161 #LBL6 | | | |
| 162 1 | | | |
| 163 5 | 85Sr | | |
| 164 3 | | | |
| 165 6 | | | |
| 166 RTN | ----- | | |
| 167 #LBL7 | | | |
| 168 1 | | | |

| LABELS | | | | | FLAGS | SET STATUS | | | | | | | | |
|--------|----------------|---|-----------|---|--------|------------|-----------|---|-------|---|--------------------------|-------------------------------------|------|------|
| A | A ₀ | B | t (dd.hh) | C | A | D | Isotope # | E | 137Cs | 0 | FLAGS | | TRIG | DISP |
| a | 51Cr | b | 57Co | c | 99mTc | d | 125I | e | 131I | 1 | ON | OFF | DEG | FIX |
| 0 | Inputs | 1 | 3H | 2 | 14C | 3 | 18F | 4 | 32P | 2 | <input type="checkbox"/> | <input checked="" type="checkbox"/> | GRAD | SCI |
| 5 | 75Se | 6 | 85Sr | 7 | 113mIn | 8 | 133Xe | 9 | 197Hg | 3 | <input type="checkbox"/> | <input checked="" type="checkbox"/> | RAD | ENG |
| | | | | | | | | | | | <input type="checkbox"/> | <input checked="" type="checkbox"/> | | n 2 |

Radioimmunoassay

| | | | | | | | | | |
|-----------|-------------------------------------|-----------|-------------------------|----|-----------------|----|-----------------|----------|-----------|
| 001 *LBLA | Initialize. | 057 F0? | B | | | | | | |
| 002 P=S | | 058 PRTX | | | | | | | |
| 003 CLRG | | 059 RCLE | | | | | | | |
| 004 P=S | | 060 - | | | | | | | |
| 005 GT02 | | 061 RCLD | | | | | | | |
| 006 *LBLB | Non-specific binding | 062 RCLE | | | | | | | |
| 007 GSD0 | counts (NSB). | 063 - | net B/B ₀ | | | | | | |
| 008 RTN | | 064 ÷ | | | | | | | |
| 009 GSB1 | | 065 F1? | | | | | | | |
| 010 ST0E | Average NSB. | 066 PRTX | | | | | | | |
| 011 RTN | | 067 F0? | | | | | | | |
| 012 *LBLC | Zero dose counts (B ₀). | 068 SPC | | | | | | | |
| 013 GSD0 | | 069 R4 | | | | | | | |
| 014 RTN | | 070 F0? | | | | | | | |
| 015 GSB1 | Average B ₀ . | 071 PRTX | Concentration. | | | | | | |
| 016 ST0D | | 072 F0? | | | | | | | |
| 017 RTN | | 073 SPC | | | | | | | |
| 018 *LBL1 | Compute average counts. | 074 LOC | log conc. (x). | | | | | | |
| 019 RCL8 | | 075 GSB3 | logit (y). | | | | | | |
| 020 RCL9 | | 076 F1? | | | | | | | |
| 021 ÷ | | 077 PRTX | Print logit. | | | | | | |
| 022 ST0I | $\bar{B} = \Sigma B/n$ | 078 XΣY | | | | | | | |
| 023 F0? | | 079 F1? | | | | | | | |
| 024 SPC | | 080 PRTX | Print log conc. | | | | | | |
| 025 F0? | | 081 F1? | | | | | | | |
| 026 PRTX | \bar{B} | 082 SPC | | | | | | | |
| 027 F0? | | 083 Σ+ | Sum x- and y-values for | | | | | | |
| 028 SPC | | 084 F0? | regression. | | | | | | |
| 029 F0? | | 085 SPC | | | | | | | |
| 030 SPC | | 086 RTN | | | | | | | |
| 031 *LBL2 | Clear for n, ΣB | 087 *LBLA | | | | | | | |
| 032 0 | | 088 RCLΣ | Compute r, m, b. | | | | | | |
| 033 ST08 | | 089 x | | | | | | | |
| 034 ST09 | | 090 P=S | | | | | | | |
| 035 R4 | | 091 RCL9 | | | | | | | |
| 036 RTN | | 092 ÷ | | | | | | | |
| 037 *LBLb | | 093 RCL8 | | | | | | | |
| 038 *LBLD | Counts for standards and | 094 XΣY | | | | | | | |
| 039 F0? | unknowns. | 095 - | | | | | | | |
| 040 PRTX | | 096 ST0B | | | | | | | |
| 041 ST+8 | | 097 ENT↑ | Σxy - (Σx Σy)/n | | | | | | |
| 042 ! | | 098 ENT↑ | | | | | | | |
| 043 ST+9 | | 099 P=S | | | | | | | |
| 044 RCL9 | | 100 S | | | | | | | |
| 045 RTN | | 101 x | | | | | | | |
| 046 *LBLE | | 102 ÷ | | | | | | | |
| 047 RCL8 | Standard concentration. | 103 P=S | | | | | | | |
| 048 RCL9 | | 104 RCL9 | | | | | | | |
| 049 ÷ | | 105 1 | | | | | | | |
| 050 ST0I | $\bar{B} = \Sigma B/n$ | 106 - | | | | | | | |
| 051 0 | | 107 ÷ | | | | | | | |
| 052 ST0B | | 108 ST0A | | | | | | | |
| 053 ST09 | | 109 PRTX | | | | | | | |
| 054 R4 | | 110 RCL8 | | | | | | | |
| 055 F0? | | 111 RCL5 | | | | | | | |
| 056 SPC | | 112 RCL4 | | | | | | | |
| REGISTERS | | | | | | | | | |
| 0 | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 |
| | | | | | | | | ΣB, used | n, used |
| S0 | S1 | S2 | S3 | S4 | S5 | S6 | S7 | S8 | S9 |
| | | | | Σx | Σx ² | Σy | Σy ² | Σxy | n |
| A | r | B | Used, m | C | b | D | B ₀ | E | NSB |
| | | | | | | | | I | \bar{B} |

Basic Statistics

| | | | | | | | | | |
|------------------|---------------------------------------|------------------|---------------------------|---------------|-----------------|---------|---------|---------|--------------------|
| 001 *LBLA | Clear Σ registers | 057 SPC | Print | | | | | | |
| 002 P Σ S | | 058 N Σ Y | | | | | | | |
| 003 0 | | 059 PRTX | f_k | | | | | | |
| 004 ST04 | | 060 X Σ Y | | | | | | | |
| 005 ST05 | | 061 PRTX | x_k ----- | | | | | | |
| 006 ST06 | | 062 *LBLI | | | | | | | |
| 007 ST07 | | 063 ENT† | | | | | | | |
| 008 ST08 | | 064 ENT† | | | | | | | |
| 009 ST09 | | 065 X | | | | | | | |
| 010 P Σ S | | 066 R† | | | | | | | |
| 011 RTN | ----- | 067 ST-9 | Σf_i | | | | | | |
| 012 *LBLB | $x_i (\Sigma +)$ | 068 X | | | | | | | |
| 013 F0? | | 069 ST-5 | $\Sigma f_i x_i^2$ | | | | | | |
| 014 PRTX | | 070 R4 | | | | | | | |
| 015 $\Sigma +$ | ----- | 071 X | | | | | | | |
| 016 RTN | | 072 ST-4 | $\Sigma f_i x_i$ | | | | | | |
| 017 *LBLb | $x_k (\Sigma -)$ | 073 I | | | | | | | |
| 018 F0? | | 074 ST-6 | | | | | | | |
| 019 SPC | | 075 RCL6 | i | | | | | | |
| 020 F0? | | 076 P Σ S | | | | | | | |
| 021 PRTX | | 077 RTN | ----- | | | | | | |
| 022 $\Sigma -$ | ----- | 078 *LBLD | Find mean and | | | | | | |
| 023 RTN | Grouped data. | 079 SPC | standard deviation. | | | | | | |
| 024 *LBLC | | 080 \bar{x} | | | | | | | |
| 025 P Σ S | | 081 PRTX | | | | | | | |
| 026 F0? | | 082 S | | | | | | | |
| 027 GT00 | | 083 PRTX | | | | | | | |
| 028 GT01 | ----- | 084 RTN | ----- | | | | | | |
| 029 *LBL0 | Print | 085 *LBLE | Find standard error and | | | | | | |
| 030 SPC | | 086 SPC | coefficient of variation. | | | | | | |
| 031 X Σ Y | | 087 S | | | | | | | |
| 032 PRTX | f_i | 088 P Σ S | | | | | | | |
| 033 X Σ Y | | 089 RCL9 | | | | | | | |
| 034 PRTX | x_i ----- | 090 P Σ S | | | | | | | |
| 035 *LBLI | | 091 IX | | | | | | | |
| 036 ENT† | | 092 \div | | | | | | | |
| 037 ENT† | | 093 PRTX | | | | | | | |
| 038 X | $x_i \quad x_i \quad x_i \quad f_i$ | 094 \bar{x} | $s_{\bar{x}}$ | | | | | | |
| 039 R† | | 095 S | | | | | | | |
| 040 ST+9 | $f_i \quad x_i^2 \quad x_i \quad f_i$ | 096 LSTX | | | | | | | |
| 041 X | Σf_i | 097 \div | | | | | | | |
| 042 ST+5 | $\Sigma f_i x_i^2$ | 098 EEX | | | | | | | |
| 043 R4 | | 099 2 | | | | | | | |
| 044 X | | 100 X | C. V. % | | | | | | |
| 045 ST+4 | $\Sigma f_i x_i$ | 101 PRTX | | | | | | | |
| 046 I | | 102 RTN | ----- | | | | | | |
| 047 ST+6 | | 103 *LBLa | Print toggle | | | | | | |
| 048 RCL6 | n | 104 F0? | | | | | | | |
| 049 P Σ S | ----- | 105 GT00 | | | | | | | |
| 050 RTN | Grouped data—correct | 106 SFP | | | | | | | |
| 051 *LBLC | error. | 107 I | | | | | | | |
| 052 P Σ S | ----- | 108 RTN | | | | | | | |
| 053 F0? | | 109 *LBL0 | | | | | | | |
| 054 GT00 | | 110 CF0 | | | | | | | |
| 055 GT01 | | 111 0 | | | | | | | |
| 056 *LBL0 | | 112 RTN | | | | | | | |
| REGISTERS | | | | | | | | | |
| 0 | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 |
| S0 | S1 | S2 | S3 | S4 Σx | S5 Σx^2 | S6 Used | S7 Used | S8 Used | S9 n, Σf_i |
| A | B | C | D | E | F | G | H | I | |

Chi-square Evaluation and Distribution

| | | | |
|----------------|--|----------------|--------------------------------------|
| 001 #LBLA | Start. | 057 RTN | |
| 002 CF1 | | 058 #LBLC | Correct erroneous $0_k \uparrow E_k$ |
| 003 CLRG | | 059 GSB7 | ($\Sigma -$). |
| 004 2 | | 060 GSB9 | |
| 005 0 | | 061 GSB7 | |
| 006 STOI | I points to R_A . | 062 STOC | |
| 007 RTN | ----- | 063 - | |
| 008 #LBLB | Input 0_i ($\Sigma +$). | 064 X^2 | |
| 009 SF1 | $F1$ set for equal E_i . | 065 RCLC | |
| 010 GSB4 | | 066 \div | |
| 011 RCLB | | 067 RCLB | |
| 012 $X \div Y$ | | 068 - | |
| 013 + | | 069 CHS | |
| 014 STOB | $\Sigma 0_i$ | 070 STOP | |
| 015 LSTX | | 071 ! | |
| 016 X^2 | | 072 $ST - i$ | |
| 017 RCLC | | 073 RCL i | |
| 018 + | | 074 RTN | |
| 019 STOC | $\Sigma 0_i^2$ | 075 #LBLD | ----- |
| 020 ! | | 076 $F1?$ | Calculate X^2 . |
| 021 $ST + i$ | | 077 GT00 | |
| 022 RCL i | i | 078 RCLB | If equal E_i , GTO 1. |
| 023 RTN | ----- | 079 GSB4 | Recall X^2 . |
| 024 #LBLB | Correct erroneous 0_k ($\Sigma -$). | 080 R/S | |
| 025 GSB7 | | 081 GT06 | |
| 026 GSB7 | | 082 #LBL0 | ----- |
| 027 GSB8 | | 083 RCLA | Calculate X^2 for equal |
| 028 GSB7 | | 084 RCLC | E_i . |
| 029 RCLB | | 085 x | |
| 030 $X \div Y$ | | 086 RCLB | |
| 031 - | | 087 \div | |
| 032 STOB | | 088 LSTX | |
| 033 LSTX | | 089 - | |
| 034 X^2 | | 090 GSB4 | X^2 |
| 035 RCLC | | 091 R/S | |
| 036 - | | 092 RCLB | Calculate E. |
| 037 CHS | | 093 RCLA | |
| 038 STOC | | 094 \div | |
| 039 ! | | 095 GSB8 | |
| 040 $ST - i$ | | 096 GSB7 | |
| 041 RCL i | | 097 RTN | |
| 042 RTN | | 098 #LBL9 | ----- |
| 043 #LBLC | Input $0_i \uparrow E_i$ ($\Sigma +$). | 099 $X \div Y$ | Print contents of Y- and |
| 044 CF1 | | 100 GSB4 | X-registers if F0 set. |
| 045 GSB9 | | 101 $X \div Y$ | |
| 046 STOC | | 102 GSB8 | |
| 047 - | | 103 RTN | |
| 048 X^2 | | 104 #LBL4 | ----- |
| 049 RCLC | | 105 GSB7 | Space and print. |
| 050 \div | | 106 #LBL8 | ----- |
| 051 RCLB | | 107 F0? | |
| 052 + | | 108 PRTX | Print. |
| 053 STOB | $\Sigma \frac{(0_i - E_i)^2}{E_i}$ | 109 RTN | |
| 054 ! | | 110 #LBL7 | ----- |
| 055 $ST + i$ | | 111 F0? | Space |
| 056 RCL i | i | 112 SPC | |

| REGISTERS | | | | | | | | | |
|-----------|--|----|---------------------|----|----|----|----|----|----|
| 0 | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 |
| S0 | S1 | S2 | S3 | S4 | S5 | S6 | S7 | S8 | S9 |
| A | B | | C | | D | | E | | I |
| n | $\Sigma 0_i, \Sigma (0_i - E_i)^2/E_i$ | | $\Sigma 0_i^2, E_i$ | | | | | | 20 |

| | | | | | |
|-----|--------|----------------------------------|-----|----------------|-------------------------------|
| 113 | RTN | ----- | 169 | e ^x | |
| 114 | *LBL E | $\nu \rightarrow \Gamma (\nu/2)$ | 170 | x | |
| 115 | GSB8 | | 171 | 2 | |
| 116 | 1 | | 172 | RCL A | |
| 117 | STO C | | 173 | Y* | |
| 118 | X=Y? | | 174 | ÷ | |
| 119 | 2 | | 175 | RCL C | |
| 120 | ÷ | | 176 | ÷ | |
| 121 | STO A | | 177 | STOE | |
| 122 | INT | | 178 | F1? | |
| 123 | LSTX | | 179 | GSB8 | |
| 124 | X≠Y? | If ν is odd, GTO 1. | 180 | F1? | |
| 125 | GT01 | | 181 | GSB7 | |
| 126 | 1 | | 182 | RTN | |
| 127 | - | | 183 | *LBL E | ----- $x \rightarrow P(x)$ |
| 128 | N! | $(\nu/2 - 1)!$ | 184 | CF1 | |
| 129 | GSB8 | | 185 | GSB5 | First find f(x). |
| 130 | GSB7 | | 186 | RCLB | |
| 131 | STOC | | 187 | RCL A | |
| 132 | R/S | ----- | 188 | ÷ | |
| 133 | *LBL1 | ν odd. | 189 | RCL E | |
| 134 | . | | 190 | x | |
| 135 | 5 | | 191 | STOE | |
| 136 | X=Y? | | 192 | 2 | |
| 137 | GT02 | | 193 | RCL A | |
| 138 | X≠Y? | | 194 | x | |
| 139 | 1 | | 195 | STO1 | |
| 140 | - | | 196 | 1 | |
| 141 | RCLC | | 197 | STOD | |
| 142 | X≠Y | | 198 | *LBL3 | ----- Sum terms of series. |
| 143 | x | | 199 | RCLB | |
| 144 | STOC | | 200 | RCL1 | |
| 145 | LSTX | | 201 | 2 | |
| 146 | GT01 | ----- | 202 | + | |
| 147 | *LBL2 | | 203 | STO1 | |
| 148 | P1 | | 204 | ÷ | |
| 149 | FX | | 205 | RCLD | |
| 150 | RCLC | | 206 | x | |
| 151 | x | | 207 | STOD | |
| 152 | STOC | | 208 | + | |
| 153 | GSB8 | | 209 | X≠Y? | |
| 154 | GSB7 | | 210 | GT03 | |
| 155 | R/S | | 211 | RCL E | |
| 156 | *LBL4 | ----- | 212 | x | |
| 157 | SF1 | $x \rightarrow f(x)$ | 213 | GSB8 | |
| 158 | *LBL5 | | 214 | RTN | |
| 159 | GSB8 | | 215 | *LBL0 | ----- |
| 160 | STOB | | 216 | F0? | Print toggle |
| 161 | RCL A | | 217 | GT00 | |
| 162 | 1 | | 218 | SF0 | |
| 163 | - | | 219 | 1 | |
| 164 | Y* | | 220 | RTN | |
| 165 | RCLB | | 221 | *LBL0 | |
| 166 | 2 | | 222 | CF0 | |
| 167 | ÷ | | 223 | 0 | |
| 168 | CHS | | 224 | RTN | |

| LABELS | | | | | FLAGS | SET STATUS | | | |
|----------|-----------------------|---------------------------------------|-----------------------|------------------------------------|---------|---|---|---|--|
| A Start | B 0 _i (Σ+) | C 0 _i ↑E _i (Σ+) | D →X ² ; E | E $\nu \rightarrow \Gamma (\nu/2)$ | 0 Print | FLAGS TRIG DISP | | | |
| 1 Print? | 0 _k (Σ-) | 0 _k ↑E _k (Σ-) | d x → f(x) | e x → P(x) | 1 Used | 0 <input type="checkbox"/> ON <input checked="" type="checkbox"/> OFF | DEG <input checked="" type="checkbox"/> | FIX <input checked="" type="checkbox"/> | |
| 0 Used | 1 Used | 2 Used | 3 Used | 4 Print, spc | 2 | 1 <input type="checkbox"/> <input checked="" type="checkbox"/> | GRAD <input type="checkbox"/> | SCI <input type="checkbox"/> | |
| 5 Used | 6 | 7 Space | 8 Print x | 9 Print x, y | 3 | 2 <input type="checkbox"/> <input checked="" type="checkbox"/> | RAD <input type="checkbox"/> | ENG <input type="checkbox"/> | |
| | | | | | | 3 <input type="checkbox"/> <input checked="" type="checkbox"/> | | n <u>2</u> | |

t Statistics

| | | | | | | | | | |
|------------------|----------------------------------|---------------------|---|---------|--------------|----------------|-----|------|----|
| 001 *LBLA | Start. | 057 \overline{fX} | | | | | | | |
| 002 0 | | 058 ST08 | s_D | | | | | | |
| 003 ST01 | | 059 RCL1 | | | | | | | |
| 004 ST02 | | 060 \overline{fX} | | | | | | | |
| 005 ST03 | | 061 \div | | | | | | | |
| 006 RTN | | 062 \div | | | | | | | |
| 007 *LBLA | Print toggle. | 063 SPC | t | | | | | | |
| 008 F0? | | 064 PRTX | | | | | | | |
| 009 GT00 | | 065 R/S | | | | | | | |
| 010 SF0 | | 066 RCLC | | | | | | | |
| 011 1 | | 067 PRTX | | | | | | | |
| 012 RTN | | 068 RCL A | | | | | | | |
| 013 *LBL0 | | 069 PRTX | | | | | | | |
| 014 CF0 | | 070 RCLB | | | | | | | |
| 015 0 | | 071 PRTX | | | | | | | |
| 016 RTN | | 072 RTN | | | | | | | |
| 017 *LBL0 | Input x_i, y_i for paired t. | 073 *LBLD | Input x_i or y_i for t for two means. | | | | | | |
| 018 F0? | | 074 GSB0 | | | | | | | |
| 019 GSB9 | | 075 ST+2 | | | | | | | |
| 020 - | | 076 X^2 | | | | | | | |
| 021 ST+2 | | 077 ST+3 | | | | | | | |
| 022 X^2 | | 078 RCL1 | | | | | | | |
| 023 ST+3 | | 079 1 | | | | | | | |
| 024 RCL1 | | 080 + | | | | | | | |
| 025 1 | | 081 ST01 | | | | | | | |
| 026 + | | 082 RTN | | | | | | | |
| 027 ST01 | | 083 *LBLd | Correct x_k or y_k for t for two means. | | | | | | |
| 028 RTN | | 084 GSB1 | | | | | | | |
| 029 *LBLb | Correct x_k, y_k for paired t. | 085 GSB0 | | | | | | | |
| 030 F0? | | 086 ST-2 | | | | | | | |
| 031 GSB0 | | 087 X^2 | | | | | | | |
| 032 - | | 088 ST-3 | | | | | | | |
| 033 ST-2 | | 089 RCL1 | | | | | | | |
| 034 X^2 | | 090 1 | | | | | | | |
| 035 ST-3 | | 091 - | | | | | | | |
| 036 RCL1 | | 092 ST01 | | | | | | | |
| 037 1 | | 093 RTN | | | | | | | |
| 038 - | | 094 *LBLc | Input d. | | | | | | |
| 039 ST01 | | 095 ST07 | | | | | | | |
| 040 RTN | | 096 RCL1 | | | | | | | |
| 041 *LBLC | Compute paired t. | 097 ST04 | Save $n_1, \Sigma x, \Sigma x^2$. | | | | | | |
| 042 RCL2 | | 098 RCL2 | | | | | | | |
| 043 RCL1 | | 099 ST05 | | | | | | | |
| 044 \div | | 100 RCL3 | | | | | | | |
| 045 ST0A | \overline{D} | 101 ST06 | | | | | | | |
| 046 RCL3 | | 102 0 | Clear for Σy . | | | | | | |
| 047 RCL2 | | 103 ST01 | | | | | | | |
| 048 X^2 | | 104 ST02 | | | | | | | |
| 049 RCL1 | | 105 ST03 | | | | | | | |
| 050 \div | | 106 RCL7 | | | | | | | |
| 051 - | | 107 GSB1 | | | | | | | |
| 052 RCL1 | | 108 GSB0 | | | | | | | |
| 053 1 | | 109 GSB1 | | | | | | | |
| 054 - | | 110 RTN | | | | | | | |
| 055 ST0C | df | 111 *LBL E | Compute t for two means. | | | | | | |
| 056 \div | | 112 RCL6 | | | | | | | |
| REGISTERS | | | | | | | | | |
| 0 | 1 n | 2 Used | 3 Used | 4 n_1 | 5 Σx | 6 Σx^2 | 7 d | 8 df | 9 |
| S0 | S1 | S2 | S3 | S4 | S5 | S6 | S7 | S8 | S9 |
| A \overline{D} | B s_D | C df | D | E | I | | | | |

t Distribution

| | | | |
|-----------------|------------------------------------|-----------------------|-------------------------------|
| 001 #LBLA | Input ν . | 057 ST03 | ----- ν odd. |
| 002 GSB5 | ----- | 058 RTN | |
| 003 ST0D | $x \rightarrow f(x)$ | 059 #LBL1 | |
| 004 RTN | | 060 . | |
| 005 #LBLB | | 061 5 | |
| 006 GSB5 | | 062 $X=Y?$ | |
| 007 ST0A | | 063 GT02 | |
| 008 RCLD | | 064 $X \neq Y$ | |
| 009 GSB6 | | 065 1 | |
| 010 ST0B | | 066 - | |
| 011 RCLD | | 067 ST43 | |
| 012 1 | | 068 GT01 | ----- |
| 013 + | | 069 #LBL2 | |
| 014 GSB2 | | 070 P i | |
| 015 ST0C | | 071 JX | |
| 016 RCLA | | 072 RCL3 | |
| 017 RCLC | | 073 x | |
| 018 RCLB | | 074 ST03 | |
| 019 + | | 075 RTN | |
| 020 P i | | 076 #LBLC | ----- $x \rightarrow P(x)$ |
| 021 RCLD | | 077 GSB5 | ----- |
| 022 x | | 078 #LBLC | Enter here from LBL D. |
| 023 JX | | 079 CF1 | |
| 024 ÷ | | 080 ST0A | |
| 025 1 | | 081 ABS | |
| 026 RCLA | | 082 RCLD | |
| 027 X^2 | | 083 ST0B | |
| 028 RCLD | | 084 RAD | |
| 029 ÷ | | 085 JX | |
| 030 + | | 086 ÷ | |
| 031 RCLD | | 087 TAN ⁻¹ | |
| 032 1 | | 088 ST02 | |
| 033 + | | 089 RCL0 | |
| 034 2 | | 090 2 | |
| 035 ÷ | | 091 ÷ | |
| 036 CHS | | 092 INT | |
| 037 Y^x | | 093 LSTY | |
| 038 x | | 094 $X \neq Y?$ | |
| 039 ST09 | | 095 GT04 | |
| 040 PRTX | | 096 0 | |
| 041 SPC | | 097 ST05 | |
| 042 RTN | | 098 #LBL6 | ----- ν even. |
| 043 #LBL0 | ----- Compute $\Gamma(\nu/2)$. | 099 RCL2 | |
| 044 1 | | 100 COS | |
| 045 ST03 | | 101 X^2 | |
| 046 $X \neq Y$ | | 102 ST03 | |
| 047 2 | | 103 RCL2 | |
| 048 ÷ | | 104 SIN | |
| 049 ST01 | | 105 ST04 | |
| 050 INT | | 106 RCL0 | |
| 051 LSTX | | 107 2 | |
| 052 $X \neq Y?$ | | 108 $X=Y?$ | |
| 053 GT01 | | 109 GT08 | |
| 054 1 | | 110 ÷ | |
| 055 - | | 111 1 | |
| 056 N! | | 112 - | |

| REGISTERS | | | | | | | | | |
|------------------|--------|------------|---------|--------|--------|--------|-----|--------|----------|
| 0 $\nu, \nu - 1$ | 1 | 2 θ | 3 Used | 4 Used | 5 Used | 6 Used | 7 R | 8 Used | 9 $f(x)$ |
| S0 | S1 | S2 | S3 | S4 | S5 | S6 | S7 | S8 | S9 |
| A x | B Used | C Used | D ν | E | F | G | H | I | J |

Notes

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Sales and service from 172 offices in 65 countries.
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00097-90106

● B C D E