

## CALCULATION OF AN ESTIMATE OF THYROXINE-BINDING GLOBULIN CAPACITY

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*An estimate of the serum thyroxine-binding globulin (TBG) may be computed from determinations of serum thyroxine and triiodothyronine uptake. A general equation for this computation is presented and a computer program for the calculation of the estimating parameters is discussed. With these methods, the regression equation for the calculated TBG and the observed TBG is the line of identity, and the correlation coefficients from determinations on data from two laboratories were +0.88 and +0.96. The calculated TBG may be used as a screening test for abnormalities of thyroxine-binding protein and as an aid in the proper interpretation of thyroid function studies.*

Alterations in the concentration of thyroxine-binding protein, especially thyroxine-binding globulin (TBG), result in abnormal values of serum thyroxine concentration. Knowledge of the TBG concentration would be of value in the clinical interpretation of abnormal serum tests of thyroid function, in the evaluation of the effects of drugs and disease states on such tests, and in the detection and elucidation of inherited TBG abnormalities (1). Direct measurements of TBG are not readily available and the test is expensive. Determinations of the triiodothyronine resin (or red cell or surface adsorbent) uptake ( $T_3U$ ) and the total serum thyroxine ( $T_4$ ), which are used to compute the free thyroxine index, are routinely available for the evaluation of thyroid function. These same two tests may also be used to estimate the capacity of thyroxine-binding protein since there is a linear inverse relationship between unbound TBG and  $T_3U$  and a direct relationship between bound TBG and  $T_4$ . Total TBG should therefore be related to a summation of some function of  $1/T_3U$  and  $T_4$ , allowing computation of an estimate of TBG from them. The

particular form that the function assumes is highly dependent on the specific method employed in the measurement of  $T_3U$ . This is true because the magnitude of change differs whereas the direction of change caused by a particular disordered state is the same among the various  $T_3U$  methods. Thus, the general equation describing TBG as a function of  $T_3U$  and  $T_4$ , where  $T_3U$  and  $T_4$  are determined by specific methods, is:

$$TBG = a \left[ \frac{1}{T_3U} \right]^m + b[T_4]^n + c. \quad (1)$$

The problem evolves into the determination of values for the coefficients and exponents,  $a$ ,  $m$ ,  $b$ ,  $n$ , and  $c$ , which yield the best estimate of TBG for the specific analytic methods employed. The purpose of this paper is to describe how this can be done.

### METHODS

$T_3U$  was performed using a surface-adsorbent technique (Tri-Tab®, Nuclear Medical Laboratories) and  $T_4$  was measured using a competitive binding assay employing a surface adsorbent (Tetra-Tab®, Nuclear Medical Laboratories). Total TBG (as maximum binding capacity of  $T_4$ ) was determined by the method of Elzinga, et al (2) by Bio-Science Laboratories. Data from two laboratories were analyzed, with our own laboratory providing one set. Our normal range of  $T_3U$  is 25–35% as we use a modification of the Tri-Tab kit in which samples are counted against a pooled normal control serum arbitrarily assigned a value of 30%. Nuclear Medical Laboratories provided the second set of data in which the normal range of  $T_3U$  using Tri-Tab is 35–45%. If the  $T_3U$  and  $T_4$  terms are expressed as

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010 DIMENSION W0(100),X(100),Y(100),AM(100),AN(100),A(100),
020&B(100),C(100),WC(100)
025 CONTINUE
030 PRINT 10
040 10 FORMAT(1X,"THIS PROGRAM EVALUATES THE EQUATION"/
050&1X,"WC=A*X**M+B*Y**N+C, WHERE X AND Y ARE EXPERIMENTALLY"/
060&1X,"OBSERVED VALUES AND A, M, B, N AND C ARE VARIABLE"/
070&1X,"COEFFICIENTS. THE COMPUTED VALUE WC IS COMPARED TO A"/
080&1X,"KNOWN VALUE W0 AND THROUGH AN ITERATIVE PROCESS THE"/
090&1X,"VARIABLES A, M, B, N, AND C ARE CHANGED IN ORDER TO"/
100&1X,"DETERMINE THE COEFFICIENT VALUES WHICH YIELD THE MIN-"/
110&1X,"IMUM DIFFERENCE BETWEEN THE COMPUTED AND OBSERVED"/
120&1X,"VALUES."////)
130 PRINT 11
140 11 FORMAT(1X,"READ IN A TWO DIGIT FIXED POINT NUMBER"/
150&1X,"CALLED NUM WHICH SPECIFIES THE NUMBER OF SETS OF"/
160&1X,"OBSERVED VALUES FOR W0, X AND Y TO BE EVALUATED."//)
170 READ 12, NUM
180 12 FORMAT(I2)
190 PRINT 13
200 13 FORMAT(1X,"READ IN THE SETS OF OBSERVED VALUES FOR"/
210&1X,"W0, X AND Y IN 3F7.3 FORMAT. EXAMPLE - "/
220&1X,"019.000001.034001.034"/)
230 DO 38 I=1, NUM
240 38 READ 1, W0(I),X(I),Y(I)
250 1 FORMAT(3F7.3)
260 PRINT 14
270 14 FORMAT(//1X,"READ IN FIVE F5.2 NUMBERS CALLED WIDTHA"/
280&1X,"WIDTHM, WIDTHB, WIDTHN, AND WIDTHC WHICH SPECIFY THE"/
290&1X,"INTERVAL WIDTHS FOR THE VARIABLE COEFFICIENTS A, M,"/
300&1X,"B, N AND C RESPECTIVELY. EXAMPLE OF AN INTERVAL WIDTH"/
310&1X,"FOR M MIGHT BE 00.25 AND FOR C, 05.00, ETC."//)
320 READ 23, WIDTHA, WIDTHM, WIDTHB,WIDTHN,WIDTHC
330 23 FORMAT(5F5.2)
340 PRINT 15
350 15 FORMAT(1X,"READ IN FIVE TWO DIGIT FIXED POINT NUMBERS"/
360&1X,"CALLED LIMA, LIMM, LIMB, LIMN AND LIMC WHICH SPECIFY"/
370&1X,"THE NUMBER OF SUCCESSIVE INTERVALS TO BE PROCESSED FOR"/
380&1X,"EACH OF THE VARIABLE COEFFICIENTS A, M, B, N, AND C"/
390&1X,"RESPECTIVELY."//)
400 READ 16, LIMA,LIMM,LIMB,LIMN,LIMC
410 16 FORMAT(5I2)
420 PRINT 17
430 17 FORMAT(1X,"READ IN FIVE F5.2 NUMBERS CALLED A(1), AM(1)"/
440&1X,"B(1), AN(1) AND C(1) WHICH SPECIFY THE INITIAL VALUES"/
450&1X,"OF EACH VARIABLE COEFFICIENT A, M, B, N AND C RESPECT-"/
460&1X,"IVELY."//)
470 READ 18, A(1),AM(1),B(1),AN(1),C(1)
480 18 FORMAT(5F5.2)
490 DO 41 I=2,LIMA
500 41 A(I)=A(I-1)+WIDTHA
510 DO 42 I=2,LIMM
520 42 AM(I)=AM(I-1)+WIDTHM
530 DO 43 I=2,LIMB
540 43 B(I)=B(I-1)+WIDTHB
550 DO 44 I=2,LIMN
560 44 AN(I)=AN(I-1)+WIDTHN
570 DO 45 I=2,LIMC
580 45 C(I)=C(I-1)+WIDTHC
590 PRINT 66
600 66 FORMAT(1X,"SUM OF DIFF"/1X,"SQUARED",7X,"A",8X,
610&"M",8X,"B",8X,"N",8X,"C"/)
620 DO 55 I=1, LIMA
630 DO 54 J=1, LIMM
640 DO 53 K=1, LIMB
650 DO 52 L=1, LIMN
660 DO 50 M=1,LIMC
670 SUMSQ =0.0
680 DO 51 N=1,NUM
690 WC(N)=A(I)*X(N)**AM(J)+B(K)*Y(N)**AN(L)+C(M)
700 51 SUMSQ=(WC(N)-W0(N))**2+SUMSQ
705 IF (SUMSQ .GT. 215.) GO TO 50
710 PRINT 99, SUMSQ, A(I), AM(J), B(K), AN(L), C(M)
720 99 FORMAT(1X, F10.3, 5(3X, F6.2))
730 50 CONTINUE
731 52 CONTINUE
732 53 CONTINUE
733 54 CONTINUE
734 55 CONTINUE
740 STOP
750 END

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**FIG. 1.** FORTRAN program to compute, by iteration of coefficients and exponents, values of TBG from patient data and to calculate sum of squares of differences between observed and calculated TBG.

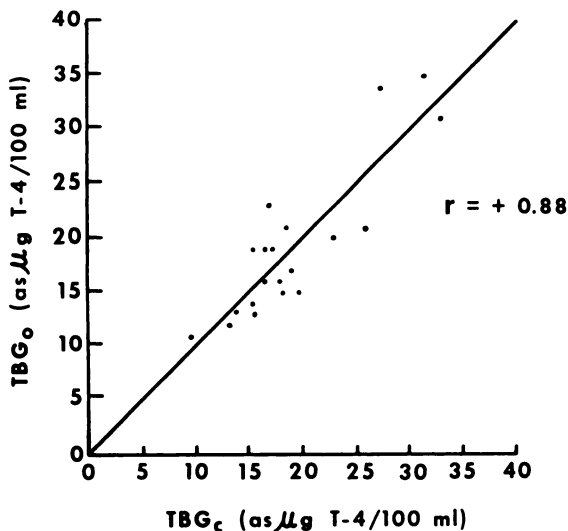


FIG. 2. Regression of calculated and observed TBG. Least-squares estimating equation is line of identity. Data used for calculating TBG were from our laboratory.

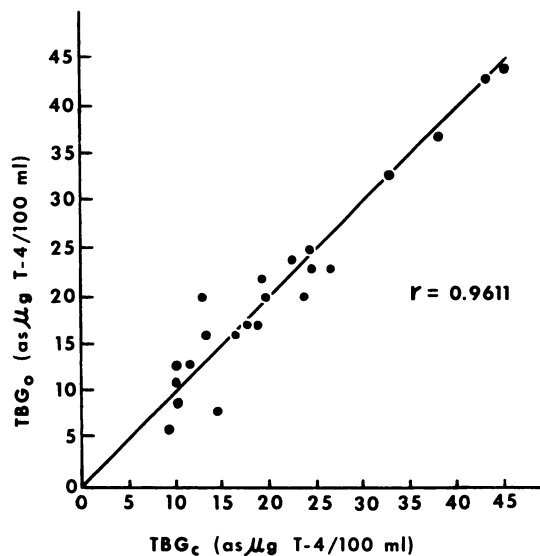


FIG. 3. Regression of calculated and observed TBG. Least-squares estimating equation is line of identity. Data used for calculating TBG were from Nuclear Medical Laboratories.

fractions of the midnormal values for the specific method (in order to increase uniformity among various assay methods), the equation becomes:

$$TBG = a \left[ \frac{T_3 U_{mid}}{T_3 U} \right]^m + b \left[ \frac{T_4}{T_4 mid} \right]^n + c. \quad (2)$$

Data from our laboratory were used to demonstrate the explicit methods employed. The  $T_3U$ ,  $T_4$ , and TBG values were obtained on serum samples of 20 patients with a variety of disorders known to affect all three tests. The variables  $a$ ,  $m$ ,  $b$ ,  $n$ , and  $c$  were initialized to some arbitrary values to compute  $TBG_c$ , a calculated TBG. The difference between the observed TBG ( $TBG_o$ ) and the  $TBG_c$  for each patient was computed, and the sum of the squares of the differences ( $TBG_c - TBG_o$ ) for each patient for the preselected values assigned to the five parameters was determined. Following an iterative scheme, the entire process was repeated for various assigned values of the parameters. The sums of squares were compared to find a minimum value and the iterative process was repeated until satisfactory convergence was obtained.

A FORTRAN computer program (Fig. 1) was employed to evaluate the equation. In the program the following symbolism was used:

$$\begin{aligned} X &= T_3 U_{mid} / T_3 U \\ Y &= T_4 / T_4 mid \\ WC &= TBG_c \\ WO &= TBG_o \end{aligned}$$

The program includes provision for the arbitrary selection of a value for the sum of squares such that only values less than this are printed. The output was inspected for the minimum value of sum of

squares and for the corresponding values of the coefficients and exponents. Iterations were then made around these values until convergence resulted.

Final determination of the coefficients  $a$ ,  $b$ , and  $c$  was made by determining the regression equation on the preliminary value of  $TBG_c$  as calculated using the values given by the iterative technique and modifying these preliminary values by the coefficients of the regression equations. Identical methods were employed in devising the regression equation for the data supplied by Nuclear Medical Laboratories.

### RESULTS

From 20 sets of results from our laboratory, and using our laboratory values of 30% for  $T_3U_{mid}$  and  $7.81 \mu\text{g}/100 \text{ ml}$  for  $T_4 mid$ , the preliminary equation determined by the iterative technique was:

$$TBG_c = 14 \left( \frac{30}{T_3 U} \right)^{1.5} + 15 \left( \frac{T_4}{7.81} \right)^{0.4} - 14.6.$$

The equation relating  $TBG_o$  and  $TBG_c$  using these parameters was:

$$TBG_o = 0.9815 (TBG_c) + 0.33.$$

Accordingly, the values for  $a$ ,  $b$ , and  $c$  were modified by multiplying by 0.9815 and adding 0.33 to the value of  $c$ . The final equation resulting was:

$$TBG_c = 13.74 \left( \frac{30}{T_3 U} \right)^{1.5} + 14.72 \left( \frac{T_4}{7.81} \right)^{0.4} - 14.00.$$

Using this equation to calculate TBG, the mean difference between the 20 pairs of values  $TBG_o$  and  $TBG_c$  was  $0.0 \pm 3.3$  (1 s.d.). Figure 2 shows the

correlation between  $TBG_o$  and  $TBG_e$  for each of the 20 patients; the correlation coefficient  $r = 0.88$ . The equation for predicting  $TBG_o$  from  $TBG_e$  was the line of identity:

$$TBG_o = 1.00 (TBG_e) + 0.00.$$

From 23 sets of data supplied by Nuclear Medical Laboratories, and using their laboratory values of 40% for  $T_3U_{mid}$  and 8.00  $\mu\text{g}/100$  ml for  $T_4_{mid}$ , the final equation resulting was:

$$TBG_e = 15.35 \left( \frac{40}{T_3U} \right)^{1.5} + 14.96 \left( \frac{T_4}{8.00} \right)^{0.5} - 9.70.$$

Using this equation to calculate TBG, the mean difference between the 23 pairs of values  $TBG_o$  and  $TBG_e$  was  $0.0 \pm 3.1$  (1 s.d.). Figure 3 shows the correlation between  $TBG_o$  and  $TBG_e$  for each of the 23 patients; the correlation coefficient  $r = 0.96$  and the equation for predicting  $TBG_o$  from  $TBG_e$  was also the line of identity.

#### DISCUSSION

The results clearly demonstrate that a relatively accurate estimate of the TBG may be computed from the  $T_3U$  and  $T_4$ .

The method employed may be adapted to the specific analytic technique used in each laboratory to compute the values of the coefficients and expo-

nents of the general equation, enabling estimations of TBG from the  $T_3U$  and  $T_4$ . While the estimated value really reflects the binding capacities of all the thyroxine-binding proteins, we have chosen to call it a TBG estimate since TBG is the binding protein of major significance. The estimated TBG is valid over a wide range of TBG values and provides a valuable tool for interpreting the serum  $T_4$  and  $T_3U$  in states in which these are affected by binding-protein changes. In addition, the TBG estimate is a useful screening test for abnormalities of TBG (or other binding proteins); thus, an abnormal TBG estimate would indicate which patients should be studied by costlier but direct assays of TBG.

The method described herein provides a means of utilizing two routine chemical determinations for calculating the concentration of a physiologically important protein, which is important to know for proper interpretation of thyroid function studies and for correct determination of the clinical state.

#### REFERENCES

1. NUSYNOWITZ ML, STRADER WJ: A new estimate of the thyroxine binding globulin capacity. *Am J Clin Pathol* 58: 718-722, 1972
2. ELZINGA KE, CARR EA, BEIERWALTES WH: Adaptation of the standard Durrum-type cell for reverse-flow paper electrophoresis. *Am J Clin Pathol* 36: 125-131, 1961

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