# COMPUTER-FOCUSING FOR AREA SCANS 

T. Nagai, T. A. linuma and S. Koda<br>National Institute of Radiological Sciences, Chiba, Japan

Scintillation area scanning is a relatively new diagnostic method in clinical medicine and its use has progressed rapidly over the years. Because the method is used to visualize the spatial distribution of radioactivity in internal organs, one would like to be able to detect and display the smallest possible lesions. Many methods using a wide range of instruments and radiopharmaceuticals have been advocated to increase the resolving power of the scanner.

At the present time, however, the resolution of area scanners is not sufficiently sharp. A multichannel focusing collimator consisting of a honeycomb of hexagonal holes is usually used, but even with this focusing collimator the region of response is broad and has a circular cross section at the focal distance. The sensitivity in scanning must also be increased, but at present this can be achieved only by sacrificing spatial resolution.

In a previous paper (1) we reported that a correction method based on iterative approximation which had been used to correct distortion in beta and gamma spectra (2) could be used to extract true information from the observed data; with this method corrected profiles obtained with a wholebody linear scanner showed a more detailed structure than did the original. We felt that a similar correction method could be used in the image of area scans.

At present a wide variety of analog techniques are used to record area-scanning data. These analog techniques, however, appear to offer less accurate recording and result in a loss of information. Moreover, they are not adaptable to computer analysis. To use digital-computer processing, it is necessary to use digital recording in which all original information in an unmodified form is collected and recorded as an array of actual numbers.

The purpose of this paper is to show how digital information suitable for computer processing can be used and how more information can be obtained from computer-corrected area scans than from the
original digital scan or conventional analog data presentation.

## METHODS

The data-collection system consisted of a commercial scanner with a $\mathrm{NaI}(\mathrm{Tl})$ crystal, 2 in . in diameter and 2 in. thick, and a 19-hexagonal-hole honeycomb collimator. Pulses from a single-channel pulse-height analyser were fed into a 128 -channel multichannel analyser used in the multiscaling mode. Because a bidimensional multiscaler was not available, the single-dimensional 128-channel multiscaler was used to present a numerical profile for each scan sweep.

One-way scanning was done at a speed of 2.7 $\mathrm{mm} / \mathrm{sec}$, and $1-\mathrm{mm}$ spacing was selected. The preset counting time in each channel of the multiscaler was 0.38 sec . Consequently each channel corresponded to the accumulation of counts from a length of 1 mm of scan sweep at the scanning speed selected. The recording of the counts for each sweep was started just when the reference point of the detector passed over the scan registration line which met the scan sweep direction at right angles in order to include precise positional information. After each scan sweep, the counts accumulated in each of the 128 channels were printed with a Hewlett Packard fast printer. During the printing the detector returned to the next starting point but spaced 1 mm perpendicular to the sweep direction. Then the multiscaler started to accumulate counts for the next scan sweep.

A channel number corresponded to the position of the detector in each scan sweep, and the number of the sweep corresponded to the position in the space direction. Thus to provide a two-dimensional array of numbers representing area scanning data,

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the sweep was repeated to cover the entire scanned area.

Optimal size of the unit area may depend upon the resolution of the collimator, and in the present study the unit area to which each count in the array corresponded was chosen to be $1 \mathrm{~mm}^{2}$.

A simple paper phantom simulating a thyroid gland was constructed to evaluate the efficiency of our method. So that the number of counts in the unit area would be as statistically meaningful as possible, the phantom contained a rather large amount of ${ }^{131}$ - $650 \mu \mathrm{c}$. Thirteen hot and cold mock tumors of various sizes were placed on the phantom. A diagram of the phantom is shown in Fig. 1 together with its autoradiogram and conventional photoscan. Items A are tumors loaded with three times the radioactivity, and items $B$ have two times the radioactivity of the remaining area. Items $C$ are cold tumors with no radioactivity. An unexpectedly high concentration of radioactivity on the perimeter was caused by a capillary phenomenon and is seen on the phantom as shown in the autoradiogram.

The phantom was placed at a focal distance of 6 cm , and the scanning was performed in air. The counts at $360 \pm 50 \mathrm{kev}$ varied from 14 to 1,336 counts, and background counts were between 0 and 3 per unit area. The array of counts obtained consisted of $126 \times 87$ elements.

Because scan data of a point source that were a fundamental measure of the resolution characteristics of the collimator were needed for a "computerfocusing" method, a ${ }^{131}$ I point source with a $2-\mathrm{mm}$ diameter was scanned in exactly the same way. From the data obtained, a $21 \times 21=441$ element array of counts was selected for computer processing.

## COMPUTER PROCESSING

In processing the digital scan obtained the computer program undertook the following steps: 1 . background subtraction, 2. smoothing the original data, 3. iterative approximation to approach true distribution and 4. again smoothing the corrected data. The recorded array of counts was punched onto computer cards and then fed into a Burroughs- 5500 digital computer. The computer time for this procedure was approximately 28 min .

Although backgre and subtraction can be preset at any level, in this case background counts in each unit area were so small that the subtraction was not performed.

Because the accumulated counts have inherent statistical variation, a data-smoothing technique is necessary, especially when the counting rate is low. This smoothing can be performed by a computer.

FIG. 1. Diagram, autoradiogram and conventional photoscan of thyroid phantom containing hot and cold mock tumors.

on collimafor used. B gives transverse response profles obtained from lines marked on A. Shaded area represents resolving power array $\|A\|$.

Each number was compared to the mean of its surrounding eight neighbors to determine whether or not it was significantly different. If it was more than one standard deviation away from the mean, it was replaced by the mean. In this way, smoothing was performed for the data of the point source and the thyroid phantom. After this initial smoothing, further computer processing was done to correct the "blur" due to lack of resolution of the collimator. Since the correction procedure has been described in a previous paper (1), only a brief explanation is given here.

For convenience, the following notations are defined;
||X|| original scanning data after smoothing, expressed in a two-dimensional array form (original image).
$X_{i j} \quad$ an element in array $\|X\|$ that corresponds to a counting rate per unit area (the ith column of the jth row); total number of elements are nm (original elements).
$\|\mathbf{A}\|$ resolving power array of a collimator that is obtained by scanning a point source of unit activity.
$\mathrm{A}_{\infty} \quad$ a central element of $\|\mathrm{A}\|$ that corresponds to a counting rate when a collimator is coaxially placed with a point source.
$A_{k l} \quad$ an element of $\|A\|$ that corresponds to a counting rate when the collimator is displaced a certain distance from the point source. $\Sigma A_{k 1}$ is normalized to unity.
The iterative approximation proceeds as follows; for simplicity calculation of an element $X_{i j}$ is shown.

In the first approximation

$$
\begin{equation*}
X_{i j}^{1}=X_{k j}+\left(X_{i j}-\sum_{k 1} A_{k 1} X_{t+k, j+1}\right) \tag{1}
\end{equation*}
$$

where $\Sigma \mathbf{A}_{\mathbf{k} 1} \mathbf{X}_{1+\mathbf{k}, j+1}$ means that the multiplication starts from $\mathrm{A}_{\infty 0} \mathrm{X}_{4 y}$; then $\mathrm{A}_{k 1}$, which corresponds to a certain distance from $A_{00}$, should be multiplied by $\mathrm{X}_{1+k, j+1}$, which corresponds to an identical distance from $X_{i y}$. The same process is repeated for all $A_{k 1}$. The calculation of Eq. 1 must be performed for all $\mathrm{X}_{1}$.
Similarly the ith approximation is

$$
\begin{equation*}
\mathbf{X}_{\mathfrak{j}}^{1}=\mathbf{X}_{1 j}^{1-1}+\left(\mathbf{X}_{1 j}-\Sigma \mathbf{A}_{\mathbf{k} 1} \mathbf{X}_{1+\mathbf{k}, j+1}^{1-1}\right) \tag{2}
\end{equation*}
$$

The approximation was stopped when $\Sigma \mathrm{A}_{\mathrm{k} 1} \mathrm{X}_{\mathrm{ij}}^{1}$ agreed with $X_{1 j}$ within the statistical standard deviation $\left(\mathrm{X}_{1 \mathrm{j}}\right)^{1 / 2}$. For all nm unit areas, the limit of approximation is (see Appendix)

$$
\begin{equation*}
\sum_{n m} \frac{\left(X_{i j}-\Sigma A_{i l} X_{1+k, j+1}^{1}\right)^{2}}{X_{i j}} \leq n m \tag{3}
\end{equation*}
$$

In the present case $\|\mathrm{X}\|$ was obtained as a 126 $\times 87$ array which corresponded to a far larger area than that of the original phantom. For $\|\mathrm{A}\|$ the number of elements was restricted to $21 \times 21=$ 441. The elements outside this array have negligible values compared to those inside, as shown in Fig. 2.
Consequently, the iterative calculation could be performed for the $\mathrm{X}_{1 j}$ which satisfied the conditions $11 \leq \mathrm{i} \leq 116$ and $11 \leq \mathrm{j} \leq 77$. Thus the number of corrected elements was $106 \times 67=7,102$, corresponding to the nm in Eq. 3.

After an iteration was completed, Eq. 3 was calculated automatically by the computer, and the calculated value was compared to 7,102 . For the present case, the values were $10,870,7,436$ and 5,998 for first, second and third iterations, respectively, and convergence of the value was quite satisfactory. According to the criterion of Eq. 3, the iteration was stopped at the third time. To prevent oscillations in the solution of the iterative approximation, smoothing was again necessary for $\mathrm{X}_{\mathrm{ij}}^{1}, \mathrm{X}_{\mathrm{ij}}^{2}$ and $X_{i j}{ }^{3}$, respectively. These final results were compared with the original elements $X_{i j}$ and the autoradiogram of the phantom.

## RESULTS

Because the immediate visual impression may be lost with fully digital display because the patterns are presented as an array of complicated numbers, some interpolation is necessary such as drawing color scans or isocount contour lines. In this case color scans and isocount contours were plotted manually from the digital display obtained to help the human visual system in the interpretation.

Figure 2 A shows a digital-to-analog converted isocount contour map of the point source. It shows that a point source appears many times larger and is shaped like a six-armed starfish, the exact shape depending on the focusing collimator used.

Figure 2 B gives the transverse response profiles
obtained from the lines marked on Fig. 2 A. The shaded areas represent the resolving power array $\|A\|$. A portion of the resolving power array normalized to unity is shown in Table 1.

Digital-to-analog converted isocontour maps of the thyroid phantom are shown in Fig. 3. Note that the difference in apparent amount of radioactivity is much greater in the computer-corrected scan (B) than in the smoothed original scan (A).

Figures $3 \mathrm{C}, \mathrm{D}$ and E show profiles obtained at the lines marked on Figs. 3 A and B . Peaks and troughs barely distinguishable on the smoothed original profiles are obvious on the corrected profiles, and the corrected curves show more of the detailed structure of the distribution. The minimum in the profile (D) represents the interlobular space, and the curve clearly shows a depression from the cold tumor in the lower pole of the left lobe.

The effect of continued iterations on profile curves obtained at the line marked $a-b$ on Figs. 3 A and $B$ can be seen in Fig. 4, which shows the first, second and third iterative approximations. This is evidence that the approximations converge satisfactorily. and the figure shows that our procedure yields a good approximation after a few iterations in this case.

Table 2 shows one of the features of the smoothed original array of the actual counts accumulated per unit area as compared to the features of the com-puter-corrected pattern shown in Table 3. Both

TABLE I. PARTIAL FEATURES OF A ARRAY NORMALIZED TO UNITY

|  |  |  |  | Space No. |  |  |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| Channel No. | 28 | 29 | 30 | 31 | 32 | 34 |  |
| 21 | 0.001018 | 0.001077 | 0.001044 | 0.001074 | 0.001041 | 0.000998 | 0.001005 |
| 22 | 0.001371 | 0.001388 | 0.001468 | 0.001421 | 0.001397 | 0.001367 | 0.001282 |
| 23 | 0.001812 | 0.001987 | 0.001923 | 0.001824 | 0.001764 | 0.001722 | 0.001421 |
| 24 | 0.002362 | 0.002419 | 0.002452 | 0.002412 | 0.002290 | 0.002131 | 0.001820 |
| 25 | 0.002919 | 0.002985 | 0.003031 | 0.002897 | 0.002839 | 0.002586 | 0.002182 |
| 26 | 0.003621 | 0.003691 | 0.003636 | 0.003639 | 0.003519 | 0.003069 | 0.002669 |
| 27 | 0.004034 | 0.004233 | 0.004340 | 0.004271 | 0.004155 | 0.003715 | 0.003252 |
| 28 | 0.004466 | 0.004777 | 0.005060 | 0.004924 | 0.004679 | 0.004202 | 0.003766 |
| 29 | 0.004777 | 0.005217 | 0.005571 | 0.005338 | 0.005052 | 0.004703 | 0.004280 |
| 30 | 0.005193 | 0.005483 | 0.005668 | 0.005538 | 0.005230 | 0.004792 | 0.004452 |
| 31 | 0.005263 | 0.005673 | 0.005743 | 0.005659 | 0.005431 | 0.004925 | 0.004412 |
| 32 | 0.005145 | 0.005477 | 0.005617 | 0.005629 | 0.005357 | 0.004906 | 0.004415 |
| 33 | 0.004856 | 0.005187 | 0.005357 | 0.005397 | 0.005132 | 0.004764 | 0.004271 |
| 34 | 0.004549 | 0.004792 | 0.005079 | 0.004904 | 0.004758 | 0.004405 | 0.003874 |
| 35 | 0.004173 | 0.004326 | 0.004484 | 0.004531 | 0.004220 | 0.003900 | 0.003389 |
| 36 | 0.003630 | 0.000808 | 0.003844 | 0.003901 | 0.003846 | 0.003666 | 0.003329 |
| 37 | 0.002938 | 0.003171 | 0.003261 | 0.003398 | 0.003305 | 0.003129 | 0.002771 |
| 38 | 0.002351 | 0.002590 | 0.002660 | 0.002739 | 0.002637 | 0.002428 | 0.002293 |
| 39 | 0.001857 | 0.002066 | 0.002103 | 0.002196 | 0.002098 | 0.002001 | 0.001913 |
| 40 | 0.001463 | 0.001591 | 0.001665 | 0.001712 | 0.001671 | 0.001569 | 0.001483 |
| 41 | 0.001161 | 0.001253 | 0.001295 | 0.001336 | 0.001311 | 0.001219 | 0.001100 |

[^1]

FIG. 3. Digital-to-analog converted isocontour maps of thyroid phantom. Difference in apparent amount of radioactivity is much greater in compufer-corrected scan (B) than in smoothed original

scan (A). C, D (above) and E (below) show profles obtained at lines marked in A and B. Peaks and troughs barely distinguishable on smoothed original profles are obvious on corrected profles.


FIG. 4. First, second and third iteretive approximations on profle curves obtained at line marked (a-b) in Fig. 3A and B.
tables correspond to the shadow area on Fig. 3 A and $B$.

In Fig. 5 digital-to-analog converted color displays of the phantom are shown, together with charts
showing the relation between counts per mit area and colors. The colors are chosen arbitrarily, each color corresponding to a given number of counts, and each element corresponding to an area of $1 \mathrm{~mm}^{2}$.
table 2. ONE OF features Of COMPUTER-SMOOTHED ARRAY OF THYROID PHANTOM*

|  | Space No. |  |  |  |  |  |  |  |  |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| Channel No. | 21 | 22 | 23 | 24 | 25 | 26 | 27 | 28 | 29 | 30 |
| 70 | 882.671 | 904.438 | 929.000 | 922.323 | 933.165 | 926.271 | 932.000 | 917.875 | 903.000 | 942.000 |
| 71 | 896.000 | 925.389 | 933.000 | 945.000 | 925.000 | 921.555 | 909.000 | 903.984 | 905.607 | 902.201 |
| 72 | 938.000 | 950.000 | 970.000 | 944.000 | 918.000 | 924.944 | 905.810 | 906.800 | 887.000 | 898.851 |
| 73 | 929.951 | 934.000 | 971.000 | 950.000 | 944.000 | 902.844 | 888.000 | 905.000 | 881.206 | 874.000 |
| 74 | 952.000 | 952.000 | 915.000 | 920.000 | 871.000 | 867.000 | 873.856 | 866.000 | 871.026 | 885.000 |
| 75 | 946.000 | 925.000 | 918.000 | 898.000 | 858.000 | 854.000 | 869.000 | 839.000 | 849.000 | 869.000 |
| 76 | 934.000 | 936.000 | 901.000 | 861.000 | 834.000 | 825.000 | 830.000 | 842.000 | 853.000 | 867.000 |
| 77 | 895.000 | 888.000 | 885.750 | 862.000 | 837.250 | 810.000 | 822.125 | 808.000 | 808.000 | 833.000 |
| 78 | 864.000 | 866.219 | 855.000 | 842.000 | 815.000 | 808.297 | 787.000 | 826.000 | 821.500 | 838.188 |
| 79 | 873.000 | 861.277 | 862.000 | 816.000 | 798.000 | 810.287 | 806.198 | 792.000 | 808.211 | 833.000 |
| 80 | 870.000 | 856.910 | 810.000 | 798.000 | 797.661 | 813.000 | 788.061 | 775.000 | 795.000 | 785.000 |
| 81 | 874.000 | 828.364 | 836.000 | 802.000 | 786.000 | 787.965 | 765.000 | 785.000 | 804.000 | 802.000 |
| 82 | 818.000 | 830.170 | 818.192 | 814.000 | 811.000 | 778.000 | 765.000 | 790.000 | 777.000 | 784.000 |
| 83 | 850.000 | 812.000 | 845.000 | 809.399 | 800.425 | 779.803 | 776.000 | 802.000 | 785.125 | 771.000 |
| 84 | 835.968 | 852.000 | 840.000 | 808.000 | 782.000 | 785.000 | 781.350 | 799.000 | 779.000 | 779.000 |
| 85 | 818.000 | 825.621 | 819.328 | 792.541 | 780.943 | 788.000 | 774.169 | 776.000 | 784.000 | 798.000 |
| 86 | 803.000 | 804.994 | 823.000 | 799.000 | 743.000 | 750.000 | 756.000 | 758.000 | 770.000 | 769.578 |
| 87 | 783.896 | 794.000 | 777.374 | 767.000 | 736.000 | 743.000 | 726.875 | 738.359 | 762.000 | 741.185 |
| 88 | 764.612 | 750.485 | 744.482 | 734.000 | 676.000 | 676.000 | 683.000 | 700.154 | 706.000 | 697.398 |
| 89 | 707.000 | 698.697 | 719.000 | 657.000 | 635.000 | 632.000 | 644.269 | 633.000 | 644.000 | 669.002 |
| 90 | 662.962 | 681.000 | 643.000 | 626.000 | 609.000 | 609.784 | 630.000 | 638.000 | 621.000 | 654.000 |

* Numbers represent the actual counts accumulated per unit area.
table 3. ONe Of features of the COMPUTER-CORRECTED ARRAY OF THYROID PHANTOM*

| Channel No. | Space No. |  |  |  |  |  |  |  |  |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  | 21 | 22 | 23 | 24 | 25 | 26 | 27 | 28 | 29 | 30 |
| 70 | 1129.557 | 1131.940 | 1112.050 | 1083.315 | 1052.715 | 1034.237 | 988.587 | 975.363 | 975.341 | 957.814 |
| 71 | 1125.139 | 1162.182 | 1129.329 | 1087.302 | 1019.217 | 987.432 | 963.503 | 907.982 | 912.491 | 897.093 |
| 72 | 1148.363 | 1164.477 | 1134.283 | 1095.165 | 1028.629 | 986.300 | 909.599 | 918.525 | 887.591 | 893.720 |
| 73 | 1176.238 | 1120.100 | 1091.584 | 1015.536 | 939.003 | 893.592 | 841.262 | 848.484 | 835.394 | 853.377 |
| 74 | 1161.186 | 1094.895 | 1026.518 | 928.387 | 856.578 | 820.003 | 796.435 | 782.432 | 816.633 | 826.096 |
| 75 | 1147.007 | 1052.280 | 983.673 | 852.338 | 772.280 | 745.640 | 732.704 | 760.075 | 804.457 | 846.137 |
| 76 | 1077.914 | 992.559 | 918.962 | 809.129 | 728.021 | 695.294 | 693.211 | 735.964 | 757.516 | 798.177 |
| 77 | 986.037 | 906.969 | 837.221 | 763.729 | 674.019 | 653.179 | 660.342 | 690.369 | 754.978 | 815.586 |
| 78 | 950.856 | 879.005 | 798.088 | 706.623 | 658.696 | 636.343 | 665.707 | 691.190 | 738.811 | 819.341 |
| 79 | 960.441 | 838.282 | 742.616 | 666.073 | 652.592 | 652.007 | 663.870 | 676.482 | 720.923 | 757.649 |
| 80 | 942.239 | 849.656 | 736.113 | 679.680 | 653.973 | 632.167 | 626.616 | 676.894 | 702.682 | 772.279 |
| 81 | 905.361 | 823.301 | 782.239 | 710.433 | 678.375 | 635.756 | 654.454 | 678.569 | 723.799 | 777.655 |
| 82 | 912.755 | 856.506 | 796.827 | 766.318 | 703.842 | 677.843 | 700.456 | 737.209 | 749.543 | 741.609 |
| 83 | 938.790 | 917.247 | 858.391 | 794.578 | 746.747 | 724.350 | 758.951 | 770.865 | 778.260 | 781.518 |
| 84 | 976.031 | 935.479 | 877.742 | 814.879 | 784.758 | 763.427 | 774.266 | 799.499 | 802.449 | 842.438 |
| 85 | 965.363 | 963.479 | 922.159 | 850.593 | 783.075 | 769.934 | 803.489 | 829.861 | 877.850 | 866.538 |
| 86 | 976.481 | 951.998 | 919.900 | 837.555 | 801.274 | 783.233 | 797.682 | 822.722 | 884.844 | 888.733 |
| 87 | 979.315 | 936.573 | 906.068 | 813.206 | 753.150 | 722.453 | 756.501 | 817.986 | 817.657 | 846.499 |
| 88 | 924.866 | 904.467 | 831.135 | 739.209 | 667.092 | 655.897 | 665.850 | 704.902 | 732.389 | 751.123 |
| 89 | 855.409 | 830.897 | 750.068 | 664.519 | 619.300 | 619.597 | 639.315 | 667.261 | 708.346 | 726.671 |
| 90 | 781.009 | 742.165 | 661.828 | 593.720 | 577.972 | 595.640 | 613.773 | 629.995 | 665.965 | 674.811 |

[^2]

FIG. 5. Digital-to-analog converted color displays of phantom are shown together with charts with relation between counts per unit area and colors. Each color corresponds to given number of counts and each element corresponds to area of $1 \mathrm{~mm}^{2}$. A shows original scan, $B$ shows smoothed scan and $C$ shows digital-fo-analog converted color scan resulting from iterative approximation.


It can be said that significantly better diagnostic information is available from the original scan (A) than the conventional photoscan. Visual impression is remarkably improved on the smoothed original scan (B), but the detailed structures are still not good enough.

Figure 5 C shows the digital-to-analog converted color scan resulting from iterative approximation. The corrected scan shows tumors strikingly clearly which can only be suspected or cannot be seen on the smoothed original scan. The three hot tumors in the right lobe are clearly separated. Note the marked increase in contrast of the cold tumor in the lower pole of the left lobe. The variations in intensity that are not observed over two small hot tumors in the central area and over the edge areas of the phantom on the smoothed original scan are obvious on the corrected scan. A small defect (e) due to an artifact is observed.

The contrast ratios, normalized to the isotope concentration of a central area (d) marked in Fig. 1, are shown in Table 4. The ratios are compared to actual values obtained by a well-type scintillation counter. This suggests that the ratios observed in the corrected data are much closer to the actual ratios than those obtained from the smoothed original data, but it seems too early to draw conclusions from this preliminary quantitative estimation.

## DISCUSSION

Generally, methods for recording scan data rely on analog means, and much of the difficulty in interpreting scans results from subjective methods. Recently, however, the availability of fast automatic digital-recording devices makes it possible to obtain numerical display of scans. Thus today, to prevent the loss of data and the introduction of a time lag by analog instruments such as a countingrate meter, conventional scanners are being replaced in some instances by actual digital scanners. Digital display provides exact data, but the vast quantities of numerical data in area scanning virtually requires the use of a digital computer.

Although the current literature (3-22) describes a number of computer techniques for scan interpretation, not enough attention has been paid to the correction for finite resolution of the collimator. Computer techniques open up the possibility of correction of this complex problem.

We have developed a new computer-processing method, "computer-focusing," to reduce the "blur" in display due to lack of resolution of the focusing
collimator and to provide increased accuracy in area scanning. In the preliminary study this method provided a satisfactorily focused display without the serious noise that is usually encountered in the Fourier transform method. Moreover, the calculation was quite simple, and the programming procedure for the computer was not very complex although time required for the computation was large.

In routine clinical applications, computation may be stopped after one or two iterative approximations because of the rapid convergence of Eq. 3. Thus the optimal compromise might be found between the resulting focused images and the cost of running the computer for each particular application. It has also been shown that the focused image may indicate not only the exact size of lesions or organs, but also the total amount of radioactivity in organs and the percent of total activity in any lesion. In the case of area scanning, however, this method is valid only under the assumption that the resolution is not affected much by the depth of the source in tissue.

In the present case, digital-to-analog converted color displays and isocount contours were drawn manually from the numerical data. Automated recording using isocontour lines or characters of increasing density, however, may possibly be accomplished by computer processing in the near future.

In principle, this computer-focusing technique can be applied to any type of scanning. one, two and three-dimensional images obtained by either moving or stationary devices. provided the data are available in digital form. However, the amount of calculation would increase enormously in the case of three-dimensional focusing.

Because the resolution depends on the collimator and the gamma-ray energy, one should make a catalog of the resolving power of a certain collimator

for various radionuclides, and then the focusing can be made for an observed scan image according to the radionuclide used.

High-resolution scans with high-energy gamma emitters, such as ${ }^{132} I^{86} \mathrm{Rb}$, ${ }^{60} \mathrm{Co}$ and ${ }^{47} \mathrm{Ca}$, etc., which have been difficult to obtain by conventional methods might be attainable with this correction technique. Moreover, the method may be used to analyze radioisotope images with smaller amounts of tracers because digital recording has a markedly greater sensitivity. Although our method requires more time at present, it may be expected that digital imaging device and on-line computer systems can be used to visualize immediately a multidimensional pattern and to perform the "computer-focusing" in a very short time.

## SUMMARY

A method for iterative approximation with a digital computer to approach the true isotope distribution in area scanning has been developed. This "computer-focusing" technique offers a means of decreasing the "blur" in area-scan images introduced by lack of resolution of the collimator, providing more accurate information.

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[^1]:    * Corresponds to a central element Aoo.

[^2]:    * Area corresponds to that shown in Table 2.

