mechanisms of tissue accumulation and washout of these compounds are not yet fully understood, evidence is emerging that their interaction with subcellular organelles and complex macromolecules might provide a new basis for advancing knowledge in important aspects of tumor biology that bear some relevance to the therapy of cancer patients.

ACKNOWLEDGMENT

The author is indebted to Dr. David Piwnica-Worms (The Mallinckrodt Institute of Radiology, Washington University, St. Louis, MO) for stimulating discussion and offering constructive criticism of this article.

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MIRDOSE: Personal Computer Software for Internal Dose Assessment in Nuclear Medicine

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Key Words: internal dosimetry; computers

J Nucl Med 1996; 37:538-546

The calculation of internal dose estimates is performed by summing the radiation absorbed in various target tissues from a

number of source organs in the body that contain significant quantities of radioactive material. In nuclear medicine, the most commonly used method for the calculation of internal dose estimates is that developed by the Medical Internal Radiation Dose (MIRD) committee, as described in various documents, but most recently summarized in the MIRD Primer (1). In this article, the expression given for the absorbed dose is:

$$D_k = A_0 \sum_j \tau_j \sum_i \frac{\Delta_i \phi(r_k \leftarrow r_j)_i}{m_k}, \qquad Eq. 1$$

Received Jul. 28, 1995; revision accepted Jul. 29, 1995.

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where D_k is the mean absorbed dose to region k (Gy); A_0 is the administered activity (Bq); τ_j is the residence time in source region j (s); Δ_i is the mean energy emitted per nuclear decay for emission type i (Gy-kg/Bq-s); $\phi_i(r_k \leftarrow r_j)$ is the fraction of energy emitted in source region j which is absorbed in target region k; and m_k is the mass of target region k (kg).

The residence time for a source region is the ratio of the cumulated activity (the total number of disintegrations) to the initial activity in the region. The units are time; typical cumulated activity units are Bq-s, and units for the initial activity are typically Bq.

One may also see the ratio of the absorbed fraction $\phi(r_k \leftarrow r_j)$ over the mass m_k given as the *specific* absorbed fraction $\Phi(r_k \leftarrow r_j)$. The quantity $\sum \Delta_i \phi_i(r_k \leftarrow r_j)/m_k$ is often referred to as the S-value (for region j irradiating region k).

A number of the parameters in this expression are rather tedious to look up and evaluate on a routine basis, and thus lend themselves well to treatment with computer programs. The summations also involve repetitive tasks that are best done by computer, leaving the analyst free to devote energy to more creative tasks. Therefore, the MIRDOSE computer software was developed several years ago (2) and has been continually supported and updated since then.

Although the software was originally described in the proceedings of a Midyear Meeting of the Health Physics Society (2), and has been distributed around the world with some limited documentation, the software has never been fully described in the open literature. Such a description serves two purposes: (a) it provides the current technical basis for the software and (b) it permits the citation of the software in an open literature reference for users who wish to use it to perform dose assessment in publications or other documents.

In addition, this article outlines some of the important differences between versions 2 and 3 of the program to help users understand variations in program output between the two versions and any effect this might have on their work.

STRUCTURE OF THE PROGRAM

Overview

The main function of the program is to provide estimates of the radiation dose per unit administered activity from user-entered source organ residence times for a given radionuclide and one or more phantoms. The program uses libraries of radionuclide decay data and specific absorbed fractions to develop S-values for the source organs chosen by the user and the target organs desired. The estimates of radiation dose per unit administered activity are given in SI and traditional units, with the two organs contributing the first and second highest percentages of the total dose; all source organ contributions to total dose may be viewed if desired. All model input and assumptions are given with the program output. The program will also provide tables of S-values for all source and target organs for a given phantom or phantoms if desired, in lieu of radiation dose estimates.

Program Data Libraries

The two major datasets needed to use Equation 1, given a set of source organ residence times, are the radionuclide decay data and the specific absorbed fractions for the various phantoms of interest. In version 2 of the program, 59 radionuclides were available (Table 1). Decay data were taken from several sources; the main source was a preliminary version of the data which was eventually published as the MIRD: Radionuclide Data and Decay Schemes (3). Some other data, however, were taken from a document by David Kocher (4) and ICRP

Publication 38 (5). These data were entered manually and proofread, and some selection was made of the number of important emissions. In version 3, all of the radionuclides in the MIRD: Radionuclide Data and Decay Schemes (3), except for those including alpha or spontaneous fission decays, were electronically transferred into the program data files (Table 2). Generally, the authors of this publication omitted transitions that did not contribute more than 0.1% to the total energy per transition in that category of emission (3).

Two major classes of emissions were considered in version 2: photons ("penetrating" emissions) and electrons and beta particles ("nonpenetrating" emissions). In addition, a third class of emission was derived, which usually constituted a minor contribution to overall dose; this category included any x- or gamma-rays with energy below 10 keV. These photons were treated as "nonpenetrating" emissions in that they were generally absorbed where they were emitted; they were given a different classification so that they would not be treated as electrons in the electron dose models for bone and marrow (described below). In version 3, this classification scheme was continued with the modification that beta particles were distinguished from monoenergetic electrons for use in the calculation of dose to small spheres, as required by the model used to calculate dose to small, unit density spheres (see Special Models And Features, below). The inclusion of alpha emitters was intentionally avoided, as alpha emitters have not been used much in nuclear medicine and because their introduction caused an increase in program complexity. The use of some alpha emitters in antibody therapy has prompted consideration of inclusion of this feature in future versions.

In both versions 2 and 3, the pediatric phantom series of Cristy and Eckerman (6) was used for the photon-specific absorbed fraction libraries, giving the user the ability to calculate dose estimates for adults (70 kg), 15-yr-olds (57 kg), 10-yr-olds (32 kg), 5-yr-olds (19 kg), 1-yr-olds (9.2 kg) and newborns (3.4 kg).

The model for the 15-yr-old has been used often as an adult female reference model. These phantoms were designed to take into account the size, shape and position of the various organs at the different ages based on available literature. The masses of these phantoms and of the organ regions in each phantom are shown in Table 3. The bone marrow in the models changes considerably with age. In the newborn, the active marrow is distributed throughout the entire length of the long bones, as well as in the other bones of the skeleton. With age, the long bones contain less and less active marrow and more and more inactive, or yellow, marrow. Thus, it is questionable whether the model for the 15-vr-old should be used for the adult female. In reviewing these data, however, it was felt that the individual variation in marrow location and mass in adult women was as great, or greater, than the difference between the 15-yr-old and the adult phantoms, and that the approximation was reasonable. Thus, it was used in this fashion for a number of years.

In version 3, the pregnant female phantom series of Stabin et al. (7) was added to the software. These phantoms were designed to represent the adult female at different stages of pregnancy. In addition, this series included a specific model for the adult female different from the 15-yr-old Cristy/Eckerman phantom. The organ masses are based on those suggested in ICRP Publication 23 (8) for the adult female. The specific absorbed fractions for the 15-yr-old were modified in the case of organ self-irradiation to account for these mass differences, while specific absorbed fractions for organ cross-irradiation were left unchanged. The bone and marrow model for the 15-yr-old was also used to represent the adult female. The

TABLE 1Radionuclides Available in MIRDOSE 2

TABLE 2Radionuclides Available in MIRDOSE 3

³ H	⁴² K	⁶⁰ Co	90Y	¹²⁷ Xe	¹⁹² ir
11C	⁴³ K	⁶⁶ Ga	⁹⁷ Ru	129	¹⁹⁵ Au
¹³ N	⁴⁵ Ca	67Ga	99mTc	130	¹⁹⁵ Hg
14C	⁵¹ Cr	⁶⁸ Ga	¹¹¹ ln	131	195mpt
¹⁵ O	⁵² Fe	⁷² Ga	^{113m} ln	¹³³ Xe	¹⁹⁷ Hg
¹⁸ F	^{52m} Mn	⁷³ Se	¹²³ mTe	¹²⁹ Cs	198Au
²² Na	⁵² Mn	⁷⁵ Se	123	¹³⁷ Cs	201TI
²⁴ Na	⁵⁷ Co	81mKr	124	¹⁵⁷ Dy	²⁰³ Hg
32P	⁵⁸ Co	⁸¹ Rb	125	¹⁶⁹ Yb	²⁰³ Pb
³⁵ S	⁵⁹ Fe	82Rb	126	178W	

masses of the adult female and the pregnant female phantoms are shown in Table 4.

Specific absorbed fractions for electrons and beta particles (plus and minus) are estimated based on the rules laid out in MIRD Pamphlet No. 11 (9) for "nonpenetrating" emissions and the masses of the target regions in the various phantoms (Tables 3 and 4). Basically, the absorbed fraction is set to 1.0 when the source and target are the same and to 0.0 when they are different, with a few exceptions. Those exceptions include:

- 1. When the source organ is the contents of a hollow organ and the target is organ wall.
- 2. When the source or the target organ is the total body.
- When the source organ is in the bone or marrow and the target organ is one in which crossfire can occur in this region.

The equations for the first two exceptions are given in MIRD Pamphlet No. 11 (9). For bone and marrow, versions 2 and 3 are completely different. In version 2, the model for bone and marrow in ICRP Publication 30 (10) was adopted, as it was felt that this was the model that would be widely used by many segments of the radiation protection community. With experience with this model and examination of its assumptions, however, it became apparent that it was considerably conservative and that the more accurate model developed by Eckerman (11) would be much more useful in nuclear medicine. A description of the new bone and marrow model is given in the article by Eckerman and Stabin (12), but a brief description and summary of some of the pertinent results will be presented (see section on Special Models and Features).

Calculational Framework

In version 2, only one phantom could be chosen at a time. In version 3, all six phantoms in the pediatric phantom series or all four phantoms in the pregnant female phantom series used with a given choice of radionuclide and set of residence times. For a given phantom, after the program receives all needed input, it proceeds to estimate the S-values it needs to calculate all of the dose estimates needed, as in Equation 1.

Starting with the first source organ, S-values are calculated for each target organ, considering each emission in the decay scheme separately. In version 2, the user was asked to specify the target organs to be studied; in version 3 the program automatically calculates dose to all 26 available target organs. If the emission is a "nonpenetrating" emission, its energy is absorbed locally as described in the rules for these emissions in MIRD Pamphlet 11 (see above).

After all radionuclide emissions are considered, the total S-value for a source/target organ combination is stored in the S-value matrix in the proper position, and the next target organ is considered. Then, the next source organ is considered until all the needed S-values are calculated.

³ H	⁶⁷ Cu	⁹⁵ Nb	¹³¹ Cs	¹⁹⁵ Au
⁷ Be	⁶² Zn	^{95m} Nb	¹³² Cs	^{195m} Au
¹¹ C	⁶⁵ Zn	⁹⁹ Mo	¹³⁴ Cs	¹⁹⁸ Au
¹⁴ C	⁶⁹ Zn	94mTc	134mCs	¹⁹⁹ Au
¹³ N	^{69m} Zn	⁹⁵ Tc	¹³⁷ Cs	¹⁹⁵ Ha
140	⁶⁶ Ga	95mTc	¹²⁸ Ba	195mHa
¹⁵ O	⁶⁷ Ga	97mTC	^{131m} Ba	¹⁹⁷ Ha
¹⁹ O	68Ga	⁹⁹ Tc	¹³³ Ba	^{197m} Ha
18F	⁷² Ga	99mTc	135mBa	²⁰³ Ha
¹⁹ Ne	⁶⁸ Ge	⁹⁷ Ru	137mpa	²⁰⁶ Ha
²² Na	72As	¹⁰³ Ru	134 a	200H
²⁴ Na	⁷³ As	103mRh	¹⁴⁰ la	²⁰¹ Tl
²⁸ Mg	⁷⁴ As	¹⁰³ Pd	¹³⁴ Ce	202TJ
28AI	⁷² Se	109Pd	¹³⁹ Ce	20 0 ∏
30p	⁷³ Se	^{109m} Ag	¹⁴¹ Ce	208]]
32p	^{73m} Se	109Cd	¹⁴⁵ Pm	210TI
33p	⁷⁵ Se	¹⁰⁹ ln	¹⁴⁷ Pm	²⁰¹ Pb
³⁵ S	^{77m} Se	111 l n	¹⁴⁵ Sm	²⁰³ Pb
³⁸ Cl	⁷⁵ Br	^{111m} ln	¹⁵³ Sm	^{204m} Pb
³⁷ Ar	⁷⁶ Br	^{113m} ln	¹⁵⁴ Eu	²¹² Pb
38K	⁷⁷ Br	¹¹⁴ ln	153Gd	²¹⁴ Pb
40K	⁸⁰ Br	^{114m} in	¹⁵⁷ Dy	²⁰⁴ Bi
42K	^{80m} Br	115m i n	159 D y	²⁰⁶ Bi
43K	825.	¹¹³ Sn	165D	Bi
150	⁸² Br	117ma	¹⁶⁵ Dy	
⁴⁵ Ca	⁷⁷ Kr	^{117m} Sn	¹⁶⁵ Er	
⁴⁷ Ca	⁷⁹ Kr	¹¹⁸ Sb	^{167m} Er	
⁴⁹ Ca	81Kr	^{118m} Sb	¹⁷¹ Er	
⁴⁶ Sc	81mKr	¹²³ Te	¹⁶⁷ Tm	
⁴⁷ Sc	83mKr	^{123m} Te	¹⁷⁰ Tm	
⁴⁹ Sc	85Kr	¹²²	¹⁷¹ Tm	
48V	85mKr	123 124	¹⁶⁹ Yb	
⁴⁸ Cr	⁷⁷ Rb	124	¹⁷⁷ Ta	
⁵¹ Cr	⁷⁹ Rb	125	¹⁷⁸ Ta	
⁵¹ Mn	⁸¹ Rb	126	¹⁷⁹ Ta	
⁵² Mn	⁸² Rb	129	¹⁸² Ta	
^{52m} Mn	^{82m} Rb	130	¹⁷⁷ W	
⁵⁴ Mn	⁸³ Rb	131	¹⁷⁸ W	
⁵² Fe	⁸⁴ Rb	132	¹⁸¹ W	
⁵⁵ Fe	⁸⁶ Rb	132mj	¹⁸⁸ W	
⁵⁹ Fe	⁸² Sr	133	¹⁸⁶ Re	
⁵⁵ Co	83Sr	¹²² Xe	¹⁸⁸ Re	
⁵⁶ Co	85Sr	123 XA	^{190m} Os	
⁵⁷ Co	^{85m} Sr	¹²⁷ Xe	¹⁹¹ Os	
⁵⁸ Co	87mSr	^{129тт} Хе	^{191m} Os	
⁶⁰ Co	⁸⁹ Sr	^{131т} Хө	¹⁹⁰ lr	
⁵⁷ Ni	90 _{Sr}	¹³³ Xe	190m1 ir	
⁶³ Ni	87 Y	133mXe	190m2 r	
⁵⁷ Cu	88 _Y	128Cs	191mlr	
⁶² Cu	90Y	¹²⁹ Cs	¹⁹² lr	
⁶⁴ Cu	⁹⁵ Zr	¹³⁰ Cs	195mpt	

After calculation of the S-values, the program checks to see if the S-values for total body as a source need to be corrected to be S-values for the "remainder of the body," i.e., the total body minus any source organs. This option is invoked if "total body" is chosen as a source along with any other organs. If "total body" is the only source chosen, or if it is not chosen at all, this option is not used.

In version 3, the name of the source organ on the input screen changes from "total body" to "remainder of the body" if any other source organs are chosen to attempt to alert the user that the residence time for the "remainder of the body" is expected for this entry. In version 2, "remainder of the body" is shown when the residence time is requested. The S-values for total body as a source are corrected according to the following formula (13):

TABLE 3Masses of Source Regions in the Cristy and Eckerman Phantom Series

			Mass (g) of organ	in each phantom		
Organ	Newborn* 3.4 kg	1 yr 9.8 kg	5 yr 19 kg	10 yr 32 kg	15 yr [†] 55–58 kg	Adult mak 70 kg
Adrenals	5.83	3.52	5.27	7.22	10.5	16.3
Brain	352	884	1260	1360	1410	1420
Breasts (including skin)	0.205	1.10	2.17	3.65	407	403
Breasts (excluding skin)	0.107	0.732	1.51	2.60	361	351
Gallbladder contents	2.12	4.81	19.7	38.5	49.0	55.7
Gallbladder wall	0.408	0.910	3.73	7.28	9.27	10.5
GI tract						
LLI contents	6.98	18.3	36.6	61.7	109	143
LLI wall	7.98	20.6	41.4	70.0	127	167
SI contents and wall	52.9	138	275	465	838	1100
stomach contents	10.6	36.2	75.1	133	195	260
stomach wall	6.41	21.8	49.1	85.1	118	158
ULI contents	11.2	28.7	57.9	97.5	176	232
ULI wall	10.5	27.8	55.2	93.4	168	220
Heart contents	36.5	72.7	134	219	347	454
Heart wall	25.4	50.6	92.8	151	241	316
Kidneys	22.9	62.9	116	173	248	299
Liver	121	292	584	887	1400	1910
Lungs	50.6	143	290	453	651	1000
Ovaries	0.328	0.714	1.73	3.13	10.5	8.71
Pancreas	2.80	10.3	23.6	30.0	64.9	94.3
Remaining tissue	2360	6400	13300	23100	40000	51800
Skeleton						
Active marrow	47	150	320	610	1050	1120
Cortical bone	0	299	875	1580	3220	4000
Trabecular bone	140	200	219	396	806	1000
Skin	118	271	538	888	2150	3010
Spleen	9.11	25.5	48.3	77.4	123	183
Testes	0.843	1.21	1.63	1.89	15.5	39.1
Thymus	11.3	22.9	29.6	31.4	28.4	20.9
Thyroid	1.29	1.78	3.45	7.93	12.4	20.7
Urinary bladder contents	12.4	32.9	64.7	103	160	211
Urinary bladder wall	2.88	7.70	14.5	23.2	35.9	47.6
Uterus	3.85	1.45	2.70	4.16	79.0	79.0
Whole body	3600	9720	19800	33200	56800	73700

^{*}Phantom and total phantom weight.

$$S(r_k \leftarrow RB) = S(r_k \leftarrow TB) \left(\frac{m_{TB}}{m_{RB}}\right) - \sum_h S(r_k \leftarrow r_h) \left(\frac{m_h}{m_{RB}}\right),$$

where $S(r_k \leftarrow RB)$ is the S-value for remainder of the body irradiating target region r_k ; $S(r_k \leftarrow TB)$ is the S-value for the total body irradiating target region r_k ; $S(r_k \leftarrow r_h)$ is the S-value for source region h irradiating target region r_k ; m_{TB} is the mass of the total body; m_{RB} is the mass of the remainder of the body, i.e., the total body minus all other source organs used in this problem; and m_h is the mass of source region h.

After all corrected S-values are available, the program simply loops over all of the source organs for each target organ, calculating the individual contributions to dose and the total dose. Each contribution to an organ's total dose is saved in a matrix so that the individual contributors may be identified in a subsequent search routine. After printing a header in which the program version is displayed, along with a notation of the user's chosen radionuclide, program output label and the date, the program proceeds one target organ at a time and prints or displays the results for each target organ as it is calculated.

In version 2, the only output option was to print to the computer's LPT1: printer port; if no printer was available, the

program would end with an error status. This limitation was removed in version 3 as the user may view the results on the screen, print them or send them to a disk file for later use. In version 3, the results are automatically shown on the screen, with options available to send the results to a file or the default printer.

With any program output in either version 2 or 3, all of the organ residence times are listed below the program output, as well as the assumptions used in either the Dynamic Bladder Model or ICRP 30 GI Tract Model (see section Special Models and Features), if used. The user also has the option, in either version, to look at all source organ contributions to a target organ's total dose; if this option is chosen, this output will be included on any program output. In version 3, the user also has the option of looking next at dose estimates for other phantoms which were initially chosen for study. If this output is to be sent to a file or printer, its output will follow that for the previous phantom; all of the above information about residence times and model assumptions will be printed again with the output.

Production of S-Value Tables

If, instead of dose estimates, the user simply wants to produce a complete table of S-values for the radionuclide and phan-

[†]Also used as adult female phantom.

TABLE 4Masses of Source Regions in the Pregnant Female Phantom Series

	Mass (g) of organ in each phantom				
Organ	Adult female (nonpregnant)	Three-month pregnant female	Six-month pregnant female	Nine-month pregnant female	
Adrenals	14	14	14	14	
Brain	1200	1200	1200	1200	
Breasts (excluding skin)	360	360	360	360	
Gallbladder contents	50	50	50	50	
Galibladder wall	8	8	8	8	
LLI contents	135	135	135	135	
LLI wali	160	160	160	160	
Small intestine contents	375	375	375	375	
Small intestine wall	600	600	600	600	
Stomach contents	230	230	230	230	
Stomach wall	140	140	140	140	
ULI contents	210	210	210	210	
ULi wali	200	200	200	200	
Heart contents	410	410	410	410	
Heart wall	240	240	240	240	
Kidneys	275	275	275	275	
Liver	1400	1400	1400	1400	
Lungs	651	651	651	651	
Ovaries	11	11	11	11	
Pancreas	85	85	85	85	
Remaining tissue*	40000	39300	41700	39500	
Skeleton					
Active marrow	1300	1300	1300	1300	
Cortical bone	3000	3000	3000	3000	
Trabecular bone	750	750	750	750	
Skin	1790	1790	1790	1790	
Spleen	150	150	150	150	
Thymus	20	20	20	20	
Thyroid	17	17	17	17	
Urinary bladder contents	160	128	107	42.3	
Urinary bladder wall	35.9	36.9	34.5	23.9	
Uterine wall	80	374	834	1095	
Fetus	_	458	1640	2960	
Placenta	_	_	310	466	
Whole body	58000	58000	61500	63700	
Whole body (maternal tissues)	56800	56400	57500	56600	

^{*}Remaining tissue is defined as the part of the phantom remaining when all defined organs have been removed. This region of the phantom has been used in the radiation transport code to model muscle for dosimetric purposes. The appropriate mass of muscle to use in such calculations in the adult female, however, is 15,500 g. The entries for this region have been rounded to two significant figures.

tom(s) chosen, that may be done by selecting this pathway. It is helpful to have all of the S-values printed on one or two pages, with columns of values for each source organ. With 24-27 source organs available, this results in the need for printing 12-14 columns per page.

Printing these results is problematic because many different printers are available. For this reason, version 2 was distributed with copies of the source code so that modifications could be made by the user to allow the S-value table to be printed in compressed print on any individual printer. In version 3, the S-values were simply printed in fewer columns, therefore using more pages and the standard printer handling routines were considered adequate. For this reason, the source code was not distributed with version 3 and is not available. Output of S-values to a file is also permitted so that the user may adapt the source output to any format using available editors.

Special Models and Features

Plot Program. Version 2 was released with a program that performed simple least-squares fitting of data to functions

involving one, two or three exponential terms. The purpose of this routine was to permit the user to calculate residence times, given a set of bioretention data. The program asked the user to enter activity for times at which measurements were taken and the corresponding values of measured activity. The program then fit the data to the desired number of exponential terms, using standard least squares methodology (14). The program returned the values of a and b for the following equation:

$$A(t) = a_1 e^{-b_1 t} + a_2 e^{-b_2 t} + \dots,$$
 Eq. 3

where A(t) is the activity at time t; a_i is the amount of activity associated with component i; and b_i is the rate coefficient for clearance of component i, (hr^{-1}) . $b_i = 0.693/T_i$, where T_i is the half-time for clearance of component i (hr).

If this expression is integrated to infinity, the residence time may be quickly estimated as:

$$\tau = \frac{a_1}{b_1} + \frac{a_2}{b_2} + \dots,$$
 Eq. 4

TABLE 5
Residence Times (HR) in the Gastrointestinal Tract at Various Ages

GI tract			Phar	ntom		
segment	Newborn	1-yr-old	5-yr-old	10-yr-old	15-yr-old	Adult*
Stomach	0.5	0.5	1.0	1.0	1.0	1.0
Small intestine	0.58	0.78	3.1	4.0	4.0	4.0
Upper large intestine	1.9	2.54	10.2	13.0	13.0	13.0
Lower large intestine	3.5	4.68	18.8	24.0	24.0	24.0

^{*}These values used for the adult and pregnant female cases.

where τ is the residence time (hr).

This expression assumes that the rate coefficient as defined above represents the sum of biological and physical elimination (the effective removal coefficient).

ICRP 30 Gastrointestinal Tract Model

In ICRP Publication 30 (10), a model for material transport through the gastrointestinal (GI) tract was proposed. In this model, material was assumed to enter the stomach and pass to the small intestine (SI), upper large intestine (ULI), lower large intestine (LLI) and be excreted in the feces at standard rates. Absorption from the SI into the blood was also allowed by the model. This model was implemented in both versions 2 and 3 to be used in cases in which activity in the GI tract was assumed to follow these kinetics.

In addition, activity was permitted to also enter the SI and pass through the rest of the system at the standard rates, to represent cases in which activity entered the GI tract from the liver. The GI transit times were based on measurements in adults and were thought to be too long to represent children. No published values of GI transit rates for children of different ages could be located. Discussions with various pediatricians, however, permitted estimation of total GI transit times in children of different ages. These total GI transit times were then scaled for each of the segments of the GI tract below the stomach in proportion to the values used for adults. The GI transit times used for the different ages are shown in Table 5.

Dynamic Bladder Model

A mathematical model which predicted the total number of disintegrations occurring in the urinary bladder, assuming input into the bladder based on a sum of exponential terms and a regular voiding pattern, was given by Cloutier et al. (15). This model was implemented in both versions of the software. The user was asked for the number of components to the input function (number of exponential terms) and for the fraction of injected activity and the biological half-time for clearance for each of the terms. The program then calculated the residence time in the bladder according to the formula given in this model.

Region-Specific Bone and Marrow Model

In version 3, the new bone and marrow model of Eckerman (11) for electron dose to bone surfaces and marrow from sources in the trabecular or cortical bone was implemented. This replaced the model in version 2, which was based on the model given in ICRP Publication 30 (10). Photon dose in both versions was based on the photon-specific absorbed fractions of Cristy and Eckerman for children and adults (6). The model used in version 3 was a significant departure from that used in

version 2 both in terms of the numerical values estimated and capability.

The ICRP 30 model for electron dosimetry of bone and marrow was based somewhat on the work of Spiers et al. (16), but was generally very conservative and had little dependence on energy. Basically, a conservatively high value for the fraction of electron energy emitted in a source absorbed in a target was chosen to represent all energies. In the Eckerman model (11), the absorbed fractions were estimated at various energies. Then, the electron and/or beta spectrum for a radio-nuclide was folded over the absorbed fraction spectrum to produce dose conversion factors. In addition, the absorbed fraction spectrum was defined for several different bone groups in the body, permitting study of the absorbed dose to bone surfaces or marrow in different regions of the skeleton.

Knowledge of the fraction of marrow existing in different bones then permitted the expression of an absorbed dose distribution in histogram format. Therefore, in version 3 this new model was implemented and users were permitted use of the most recent photon- and electron-absorbed fractions. Users were also given the ability to look at marrow dose distributions and dose-volume histograms. An example of the output from the program showing this information is given in Figure 1. The residence times used to generate these results are those shown in Table 9.

Absorbed Dose to Small, Unit Density Spheres

Calculation of the absorbed dose to small, unit density spheres is often of interest, e.g., in the dose calculation to a tumor or to a small organ. Absorbed fractions for photons, given the assumption that the activity is distributed uniformly throughout the sphere, have been available for a number of years (17,18). Absorbed fractions for electrons and beta particles were recently published for a number of sphere sizes and emission energies (19). These datasets were combined in version 3 to permit the user to calculate S-values for any radionuclide in the program very easily (Fig. 2). An example of the output of this module is shown in Table 6.

RESULTS AND DISCUSSION

Tables 7–9 show a comparison of results generated by versions 2 and 3 for the reference adult phantom (70 kg), given a set of entered residence times. In Table 7, values for ¹⁴C-glucose are shown. Carbon-14 is a pure beta emitter, and the model shown for ¹⁴C-labeled glucose uses only four organs: the brain, liver, urinary bladder contents and remainder of the body.

The results between the two versions are virtually identical, with the exception of the dose to the bone marrow and bone surfaces. This is *not* due to the change in the bone and marrow model, but in the way that the dose from the remainder of the body to these organs is estimated.

In version 2, uniform activity in the remainder of the body was assumed to be uniform in marrow and bone and the absorbed fractions for electrons irradiating marrow and bone surfaces were applied to estimate S-values. In version 3, it was assumed that all organs should receive the same dose from activity uniform in the remainder of the body. This was the method employed in MIRD Pamphlet No. 11, and is a more reasonable approach.

Table 8 shows a comparison of results from the two codes for ^{99m}Tc-labeled MIBI (methoxyisobutyl isonitrile). The results are quite similar for all organs in this case.

Table 9 shows a comparison of results for a monoclonal antibody labeled with ¹³¹I, which has some uptake in the red marrow and the bone. In this table, the differences between the

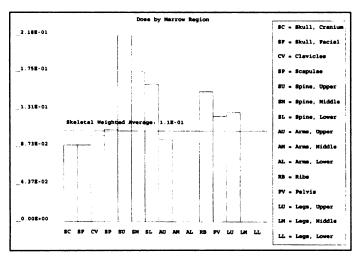


FIGURE 1. Sample of marrow dose report available in MIRDOSE versions 3.0 and 3.1. This plot shows marrow dose to different regions of the skeleton. On the computer screen, the histogram regions are shaded in different colors and on the printed output they are shown in different shades of grey (on a black and white printer) or color (on a color printer). For purposes of reproduction here, they are left unshaded.

uptake in the bone and marrow dose are clearly seen. All of the reasons for these differences will not be immediately clear from this example. The reader is directed to the paper by Eckerman and Stabin (12) for a more complete explanation of the new bone model and its differences from previous models.

Some minor differences may be noted for many other organs between the two codes. This may be attributed to a difference in

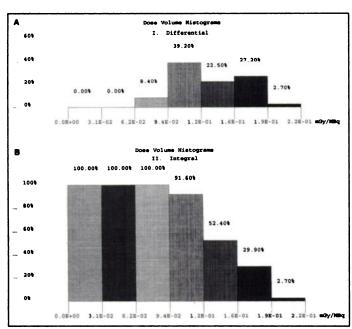


FIGURE 2. (A) Sample of the differential marrow dose volume histogram report available in MIRDOSE versions 3.0 and 3.1. This figure shows the fractions of marrow receiving absorbed doses between the values on the abscissa, based on the data in Figure 1 and assumed fractions of marrow in the different regions of the skeleton. On the computer screen, the histogram regions are shaded in different colors, and on the printed output, they are shown in different shades of grey (on a black and white printer) or color (on a color printer). (B) Sample of the integral marrow dose volume histogram report available in MIRDOSE version 3.0 and 3.1. This figure shows the fractions of marrow receiving absorbed doses greater than or equal to the values on the abscissa, based on the data in Figure 1 and assumed fractions of marrow in the different regions of the skeleton. On the computer screen, the histogram regions are shaded in different colors and on the printed output they are shown in different shades of grey (on a black and white printer) or color (on a color printer).

the marrow mass between the two codes. MIRDOSE 2 uses a marrow mass of 1500 g, one which has been widely used and quoted.

MIRDOSE 3 uses a marrow mass of 1120 g, which was actually the original mass of active marrow suggested in *ICRP Publication 23* (8), whereas the value of 1500 g is the mass of marrow which contains some red marrow (but which actually also contains some yellow marrow). When this slight mass difference is introduced into the calculation for the remainder of the body S-value correction (Equation 2), the S-values for the remainder of the body to the different organs of the body become slightly higher, and thus the doses are increased slightly.

Effective Dose Equivalent and Effective Dose

In version 3, the quantities effective dose equivalent (10) and effective dose (20) were also calculated and added to the list of dose estimates given by the program. In theory, these quantities permit the representation of a nonuniform internal dose as a single value, which is the dose equivalent to that which the whole body could uniformly receive that would result in the same overall risk as the actual nonuniform dose distribution received. This may permit comparison of the radiation risk of different diagnostic agents (e.g., a 99mTc-labeled heart agent compared to 201Tl) using a single number, or comparison of nuclear medicine and x-ray procedures, etc.

The two quantities—effective dose equivalent and effective dose—are identical in concept. They are different in name, in the numerical values of the organ risk weighting factors assigned (Table 10) and slightly different in the scheme used to estimate dose to remainder organs. Both values are given in version 3 for comparison and selection as the user chooses.

These quantities were originally designed for use in radiation protection programs (10), but their use has been suggested for nuclear medicine (21-23) by the ICRP (who designed it) and

TABLE 6
S-Values for Self-Irradiation: Small Unit Density Spheres, lodine-131

	Self-dose	S-value
Sphere mass (g)	(mGy/MBq-s)	(rad/μCi-hr)
0.01	2.34E+00*	3.11E+01*
0.10	2.70E-01*	3.60E+00*
0.50	5.70E-02*	7.59E-01*
1.00	2.95E-02	3.93E-01
2.00	1.50E-02	2.00E-01
4.00	7.64E-03	1.02E-01
6.00	5.16E-03	6.87E-02
8.00	3.89E-03	5.18E-02
10.00	3.13E-03	4.17E-02
20.00	1.58E-03	2.11E-02
40.00	8.10E-04	1.08E-02
60.00	5.55E-04	7.39E-03
80.00	4.26E-04	5.68E-03
100.00	3.45E-04	4.59E-03
300.00	1.21E-04	1.61E-03
400.00	9.23E-05	1.23E-03
500.00	7.49E-05	9.97E-04
600.00	6.30E-05	8.39E-04
1000.00	3.90E-05	5.20E-04
2000.00	2.05E-05	2.73E-04
3000.00	1.41E-05	1.88E-04
4000.00	1.09E-05	1.45E-04
5000.00	8.89E-06	1.18E-04
6000.00	7.55E-06	1.01E-04

*Electron/beta only.

TABLE 7Comparison of MIRDOSE Versions 2 and 3 Results: Pure Beta Emitter in Four Organs: Carbon-14-Labeled Glucose

TABLE 8
Comparison of MIRDOSE Versions 2 and 3 Results: Photon and Electron Emitter in Several Organs: Technetium-99m-MIBI*

	Estimated radiation dose (mGy/MBq)		
Target organ	MIRDOSE 2	MIRDOSE 3	
Adrenals	2.47E-02	2.47E-02	
Brain	3.17E-02	3.17E-02	
Breasts	2.47E-02	2.47E-02	
Gallbladder wall	2.47E-02	2.47E-02	
LLI wall	2.47E-02	2.47E-02	
Small intestine	2.47E-02	2.47E-02	
Stomach	2.47E-02	2.47E-02	
ULI wali	2.47E-02	2.47E-02	
Heart wall	2.47E-02	2.47E-02	
Kidneys	2.47E-02	2.47E-02	
Liver	4.91E-02	4.91E-02	
Lungs	2.47E-02	2.47E-02	
Muscle	2.47E-02	2.47E-02	
Ovaries	2.47E-02	2.47E-02	
Pancreas	2.47E-02	2.47E-02	
Red marrow	3.24E-02	2.47E-02	
Bone surfaces	2.21E-02	2.47E-02	
Skin	2.47E-02	2.47E-02	
Spleen	2.47E-02	2.47E-02	
Testes	2.47E-02	2.47E-02	
Thymus	2.47E-02	2.47E-02	
Thyroid	2.47E-02	2.47E-02	
Urinary bladder wall	5.45E-02	5.46E-02	
Uterus	2.47E-02	2.47E-02	
Total body	2.55E-02	2.55E-02	
Effective dose equivalent	2.93E-02*	2.84E-02	
Residence times			
Brain	1.58E+00 hr		
Liver	3.29E+00 hr		
Urinary bladder contents	6.25E-01 hr		
Remainder of the body	6.08E+01 hr		

*Effective dose equivalent not	available in version 2,	but calculated based
on the estimated organ doses.		

others. There is some controversy about the use of these quantities at the time of this writing (24,25), but the quantities are calculated in MIRDOSE 3 and offered to the user, in case they are of interest.

CONCLUSION

The MIRDOSE computer software greatly facilitates the calculation of internal radiation dose estimates by the MIRD technique. The user need only calculate organ residence times, enter them into the program and radiation dose estimates for all organs are estimated, including the effective dose equivalent and effective dose (in version 3). The program makes use of standard and most up-to-date models used in internal dosimetry. This results in standardization of dose estimates calculated from a given set of residence times, and should greatly enhance the ability of users, manufacturers, regulators and others to interpret radiation dose estimates.

Finally, it should be made clear that this program is in no way associated with the MIRD Committee of the Society of Nuclear Medicine. The name of the program implies only that it uses the MIRD technique. The MIRD Committee wishes it to be known that it does not endorse the MIRDOSE program, its input data, methods or results.

	Estimated radiation dose (mGy/MBq)		
Target organ	MIRDOSE 2	MIRDOSE 3	
Adrenals	6.26E-03	6.25E-03	
Brain	1.83E-03	1.83E-03	
Breasts	1.85E-03	1.85E-03	
Gallbladder wall	2.96E-02	2.96E-02	
LLI wall	4.19E-01	4.20E-01	
Small intestine	7.11E-02	7.09E-02	
Stomach	1.28E-02	1.28E-02	
ULI wali	1.70E-01	1.70E-01	
Heart wall	4.96E-03	4.95E-03	
Kidneys	2.31E-02	2.31E-02	
Liver	8.21E-03	8.19E-03	
Lungs	2.75E-03	2.75E-03	
Muscle	8.82E-03	8.81E-03	
Ovaries	6.25E-02	6.24E-02	
Pancreas	8.78E-03	8.77E-03	
Red marrow	1.25E-02	1.24E-02	
Bone surfaces	1.30E-02	1.31E-02	
Skin	3.40E-03	3.39E-03	
Spleen	8.63E-03	8.62E-03	
Testes	7.91E-03	7.90E-03	
Thymus	2.46E-03	2.46E-03	
Thyroid	2.22E-03	2.22E-03	
Urinary bladder wall	5.37E-02	5.37E-02	
Uterus	3.39E-02	3.39E-02	
Total body	1.08E-02	1.07E-02	
Effective dose equivalent	6.31E-02 [†]	6.31E-02	
Residence times			
Gallbladder contents	7.35E-02 hr		
Lower large intestine contents	8.86E+00 hr		
Small intestine contents	1.48E+00 hr		
Upper large intestine contents	4.80E+00 hr		
Heart wall	5.30E-02 hr		
Kidneys	2.90E-01 hr		

Urinary bladder

Remainder of the

contents

body

Liver

Lungs

Spleen

1.86E-01 hr

4.90E-02 hr

2.40E-02 hr

7.80E-01 hr

3.86E+00 hr

ACKNOWLEDGMENTS

This work was performed for the U.S. Department of Energy under contract DE-AC05-76OR00033 and for the U.S. Food and Drug Administration under Interagency Agreement No. FDA 224-75-3016, DOE 40-286-71. This article has been authored by a contractor of the U.S. Government under contract DE-AC05-76OR00033. Accordingly, the U.S. Government retains a nonexclusive, royalty-free license to publish or reproduce the published form of the contribution, or allow others to do so, for U.S. Government purposes.

^{*}Methoxyisobutyl isonitrile.

[†]Effective dose equivalent not available in version 2, but calculated based on the estimated organ doses.

TABLE 9

Comparison of MIRDOSE Versions 2 and 3 Results: Photon and Electron Emitter in Several Organs, Including the Red Marrow and Bone: Iodine-131-Labeled Monoclonal Antibody

	Estimated radiation dose (mGy/MBq)		
Target organ	MIRDOSE 2	MIRDOSE 3	
Adrenals	4.16E-02	4.19E-02	
Brain	9.25E-03	9.47E-03	
Breasts	8.77E-03	9.16E-03	
Galibladder wali	2.78E-02	2.83E-02	
LLI wall	1.62E-02	1.67E-02	
Small intestine	2.05E-02	2.10E-02	
Stomach	1.95E-02	2.00E-02	
ULI wall	1.98E-02	2.02E-02	
Heart wall	1.38E-02	1.43E-02	
Kidneys	1.57E+00	1.56E+00	
Liver	4.68E-02	4.68E-02	
Lungs	1.29E-02	1.32E-02	
Muscle	1.38E-02	1.42E-02	
Ovaries	1.69E-02	1.74E-02	
Pancreas	3.15E-02	3.19E-02	
Red marrow	1.92E-01	1.07E-01	
Bone surfaces	2.65E-01	9.93E-02	
Skin	9.40E-03	9.72E-03	
Spleen	5.87E-02	5.88E-02	
Testes	9.63E-03	1.01E-02	
Thymus	1.02E-02	1.07E-02	
Thyroid	9.58E-03	9.95E-03	
Urinary bladder wall	1.34E-01	1.34E-01	
Uterus	1.83E-02	1.88E-02	
Total body	2.51E-02	2.51E-02	
Effective dose equivalent	1.49E-01*	1.34E-01	

Kidneys	3.71E+00 hr
Liver	4.17E-01 hr
Red marrow	1.50E+00 hr
Cortical bone	5.00E-01 hr
Trabecular bone	5.00E-01 hr
Spleen	4.17E-02 hr
Urinary bladder contents	4.00E-01 hr
Remainder of the body	2.60E+00 hr

*Effective dose equivalent not available in version 2, but calculated based on the estimated organ doses.

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TABLE 10

Comparison of Organ Risk Weighting Factors for Effective Dose

Equivalent and Effective Dose

Organ	Effective dose equivalent (ICRP 30)	Effective dose (ICRP 60)
Gonads	0.25	0.20
Red marrow	0.12	0.12
Colon		0.12
Lungs	0.12	0.12
Stomach	_	0.12
Bladder	-	0.05
Breasts	0.15	0.05
Liver	-	0.05
Esophagus	_	0.05
Thyroid	0.03	0.05
Skin		0.01
Bone surfaces	0.03	0.01
Remainder	0.30*	0.05 [†]

"The weighting factor for the remainder in the effective dose equivalent is to be divided equally among the five highest organs not explicitly mentioned in this list which receive the highest absorbed doses.

[†]The weighting factor for the remainder in the effective dose is to be divided equally among ten organs not explicitly mentioned in this list but which were assigned by the ICRP (adrenals, brain, ULI, small intestine, kidneys, muscle, pancreas, spleen, thymus and uterus).

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