

MASTER

**Timed: A Computer Program for
Calculating Cumulated Activity of a
Radionuclide in the Organs of the
Human Body at a Given Time,
 t , After Deposition**

S. B. Watson
W. S. Snyder
M. R. Ford

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Printed in the United States of America. Available from
National Technical Information Service
U.S. Department of Commerce
5285 Port Royal Road, Springfield, Virginia 22161
Price: Printed Copy \$6.75; Microfiche \$3.00

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Contract No. W-7405-eng-26

COMPUTER SCIENCES DIVISION

TIMED: A COMPUTER PROGRAM FOR CALCULATING CUMULATED ACTIVITY
OF A RADIONUCLIDE IN THE ORGANS OF THE HUMAN BODY
AT A GIVEN TIME, t , AFTER DEPOSITION

S. B. Watson
Computer Sciences Division

W. S. Snyder and M. R. Ford
Health Physics Division

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Date Published: December 1976

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ACKNOWLEDGMENTS

The authors wish to express their thanks to Dr. D. E. Arnurius for his assistance in modifying the GEAR computer package for use with our particular problem.

B. D. McCarter deserves a special thanks for the preparation of the manuscript.

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ABSTRACT

TIMED is a computer program designed to calculate cumulated activity, $\Sigma Y_i(t)$, in the various source organs, Y_i , at some time t after deposition. TIMED embodies a system of differential equations which describes activity transfer in the lungs, gastrointestinal tract, and other organs of the body. This system accounts for delay of transfer of activity between compartments of the body and radioactive daughters.

The computer program contains routines which are written in either IBM System/360 or System/370 FORTRAN or IBM System/360 Assembler language. The code is executable on the IBM System/360 or System/370 machines and requires a minimum of 310 K core storage for execution.

I. INTRODUCTION

Average dose equivalent in an organ T of the body from an internally deposited emitter is given by

$$DE(T, t) = \sum_i U_{Y_i}(t) * S(T \rightarrow Y_i) \quad (1.1)$$

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where $U_{Y_i}(t)$ represents the cumulated activity measured in microcurie-days ($\mu\text{Ci-days}$) in the various source organs, Y_i , at some time t after deposition, and $S(T+Y_i)$ represents dose to a target organ T due to 1 $\mu\text{Ci-day}$ of the emitter in the source.¹ S has been tabulated for about 160 radionuclides^{2,3} and for various source and target organs, but with the proviso that the activity is always considered to be uniformly distributed in the source organ. This paper discusses the computer code, TIMED, which calculates the cumulated activity, $U_{Y_i}(t)$, per microcurie intake at some time t after intake.

In principle, this calculation is simple in that if $f_{Y_i}(t)$ represents the fraction of activity in microcuries (μCi), which entered the body at time 0 that is present in an organ Y_i at time t , the cumulated activity in that organ from time 0 to time t_1 , is given by

$$U_{Y_i}(t_1) = \int_0^{t_1} f_{Y_i}(t) dt \quad (\mu\text{Ci-days}) . \quad (1.2)$$

However, the calculation of $U_{Y_i}(t)$ is somewhat difficult due to the complex nature of the transfer rates between sections of the body and the possible presence of daughter radionuclides.

This paper discusses the model of activity transfer and the computational procedures that are used in the code, TIMED. Also presented are a description of the program and detailed user information.

II. DESCRIPTION OF THE MODEL

The model of activity transfer in the body is based on the concept of a system of compartments with constant rates of transfer to other compartments or out of the body. These compartments may be an organ or some subsystem of the body (lungs, gastrointestinal tract, blood, etc.). The system must be general enough to allow for delay of transfer of some of the activity and must also account for radioactive daughters. This system has been adequately represented by a set of differential equations which are described below for the lungs, gastrointestinal tract, and other organs. The following is taken in part from Ref. 1 which should be consulted for the detailed explanation. Many of the errors present in Ref. 1 are corrected in this discussion.

Lungs

The lung model, as described in the report of the ICRP Task Group on Lung Dynamics,⁴ and as revised in ICRP Publication 19,⁵ consists of a nasal-pharyngeal region (N-P), a tracheo-bronchial region (T-B), a pulmonary region (P), and the lymph nodes (L). Deposition is governed by the activity median aerodynamic diameter (AMAD) of the aerosol (see Ref. 4). The pathways of transfer and the deposition and transfer constants are shown in Fig. 2.1.

The differential equations appropriate for this model are given below. Taking $X_a^n, X_b^n, X_c^n, X_d^n, X_e^n, X_f^n, X_g^n, X_h^n, X_i^n, X_j^n, X_k^n, X_l^n$ as the activities of the parent ($n=1$) and daughters ($n=2, 3, \dots$) in a compartment, moving by the pathway with the appropriate subscript, and

COMPARTMENT		CLASS					
		D		W		Y	
		T	F	T	F	T	F
N-P ($D_{N-P}=0.30$)	a	0.01	0.5	0.01	0.1	0.01	0.01
	b	0.01	0.5	0.40	0.9	0.40	0.99
T-B ($D_{T-B}=0.08$)	c	0.01	0.95	0.01	0.5	0.01	0.01
	d	0.2	0.05	0.2	0.5	0.2	0.99
P ($D_P=0.25$)	e	0.5	0.8	50	0.15	500	0.05
	f	n.a.	n.a.	1.0	0.4	1.0	0.4
	g	n.a.	n.a.	50	0.4	500	0.4
	h	0.5	0.2	50	0.05	500	0.15
L	i	0.5	1.0	50	1.0	1000	0.9

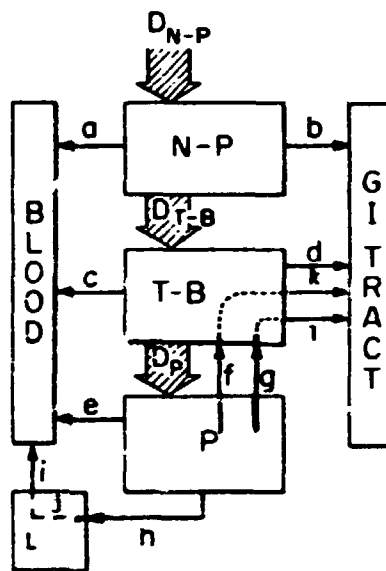


Fig. 2.1. Values for the Removal Half Times, T_a Through T_i , and Compartmental Fractions, F_a Through F_i , Are Given for Each of the Three Classes of Retained Materials. The values given for D_{N-P} , D_{T-B} , and D_P (left column) are based on an aerosol with an AMAD of $1 \mu\text{m}$.⁴ The schematic drawing identifies the various clearance pathways in the model, a through i , in relation to the initial depositions D_{N-P} , D_{T-B} , and D_P and the three respiratory regions, $N-P$, $T-B$, and P . The lymphatic clearance for class Y compounds indicates that a 90% regional fraction follows a 1000-day biological half time. The remaining 10% is presumed to be permanently retained in the nodes and is subject only to radioactive decay.

using \dot{X} to represent the derivative with respect to time ($X \equiv X(t)$ and $\dot{X} \equiv \dot{X}(t) \equiv \frac{dX(t)}{dt}$), the equations for the lungs are the following:

$$\begin{aligned}
 \dot{X}_a^{n+1} &= -X_a^{n+1}(\lambda_a^{n+1} + \lambda_r^{n+1}) + (X_a^n + X_b^n)\lambda_r^{n+1}F_a^{n+1}, \\
 \dot{X}_b^{n+1} &= -X_b^{n+1}(\lambda_b^{n+1} + \lambda_r^{n+1}) + (X_a^n + X_b^n)\lambda_r^{n+1}F_b^{n+1}, \\
 \dot{X}_c^{n+1} &= -X_c^{n+1}(\lambda_c^{n+1} + \lambda_r^{n+1}) + (X_c^n + X_d^n)\lambda_r^{n+1}F_c^{n+1}, \\
 \dot{X}_d^{n+1} &= -X_d^{n+1}(\lambda_d^{n+1} + \lambda_r^{n+1}) + (X_c^n + X_d^n)\lambda_r^{n+1}F_d^{n+1}, \\
 \dot{X}_e^{n+1} &= -X_e^{n+1}(\lambda_e^{n+1} + \lambda_r^{n+1}) + (X_e^n + X_f^n + X_g^n + X_h^n)\lambda_r^{n+1}F_e^{n+1}, \\
 \dot{X}_f^{n+1} &= -X_f^{n+1}(\lambda_f^{n+1} + \lambda_r^{n+1}) + (X_e^n + X_f^n + X_g^n + X_h^n)\lambda_r^{n+1}F_f^{n+1}, \\
 \dot{X}_g^{n+1} &= -X_g^{n+1}(\lambda_g^{n+1} + \lambda_r^{n+1}) + (X_e^n + X_f^n + X_g^n + X_h^n)\lambda_r^{n+1}F_g^{n+1}, \\
 \dot{X}_h^{n+1} &= -X_h^{n+1}(\lambda_h^{n+1} + \lambda_r^{n+1}) + (X_e^n + X_f^n + X_g^n + X_h^n)\lambda_r^{n+1}F_h^{n+1}, \\
 \dot{X}_i^{n+1} &= -X_i^{n+1}(\lambda_i^{n+1} + \lambda_r^{n+1}) + X_h^{n+1}\lambda_h^{n+1}F_i^{n+1} + (X_i^n + X_j^n)\lambda_r^{n+1}F_i^{n+1}, \\
 \dot{X}_j^{n+1} &= -X_j^{n+1}\lambda_r^{n+1} + X_h^{n+1}\lambda_h^{n+1}(1 - F_i^{n+1}) + (X_i^n + X_j^n)\lambda_r^{n+1}(1 - F_i^{n+1}), \\
 \dot{X}_k^{n+1} &= -X_k^{n+1}(\lambda_d^{n+1} + \lambda_r^{n+1}) + X_k^n\lambda_r^{n+1} + X_f^{n+1}\lambda_f^{n+1}, \\
 \dot{X}_l^{n+1} &= -X_l^{n+1}(\lambda_d^{n+1} + \lambda_r^{n+1}) + X_l^n\lambda_r^{n+1} + X_g^{n+1}\lambda_g^{n+1}.
 \end{aligned} \tag{2.1}$$

F_ξ^{n+1} is the compartmental fraction from Fig. 2.1 for the $n+1$ st radio-nuclide of the chain for pathway ξ . The quantity λ_ξ^{n+1} is $\ln 2/T_\xi^{n+1}$

where T_{ξ}^{n+1} is the removal half time in days taken from Fig. 2.1 for the $n+1$ st radionuclide of the chain for pathway ξ . Similarly, λ_r^{n+1} is $\ln 2 / T_r^{n+1}$ where T_r^{n+1} is the radioactive half life of the $n+1$ st daughter.

The terms with negative signs correspond to elimination by biological transfer or by radioactive decay. The terms involving a superscript n refer to activity in microcuries of the preceding radionuclide of the chain which produces $\lambda_r^{n+1} \mu\text{Ci}$ of the following radionuclide per unit time, and these are redistributed according to the values of F for the radionuclide.

For each chain of radionuclides there are 12 differential equations corresponding to the lung model for each radionuclide. The equations are valid for the parent radionuclide, that is $n=0$, if $x_{\xi}^0=0$ for all values of t and all subscripts ξ , thus greatly simplifying the differential equations for the parent.

The initial conditions used are based on the assumption that 1 μCi of an aerosol of AMAD=1 has been inhaled, although other starting values of $D_3 \equiv D_{N-P}$, $D_4 \equiv D_{T-B}$, and $D_5 \equiv D_P$ may be used (see Ref. 4).

$$\begin{aligned}
 x_a^1(0) &= D_3 F_a^1, & x_e^1(0) &= D_5 F_e^1, \\
 x_b^1(0) &= D_3 F_b^1, & x_f^1(0) &= D_5 F_f^1, \\
 x_c^1(0) &= D_4 F_c^1, & x_g^1(0) &= D_5 F_g^1, \\
 x_d^1(0) &= D_4 F_d^1, & x_h^1(0) &= D_5 F_h^1,
 \end{aligned} \tag{2.2}$$

$$x_i^1(0) = x_j^1(0) = x_k^1(0) = x_l^1(0) = 0,$$

$$x_{\xi}^n(0) = 0 \text{ for } n > 1 \text{ and for all subscripts } \xi.$$

Upon solution of the differential equations for the X 's which are given as activity in μCi 's, one calculates the cumulated activity ($\mu\text{Ci-days}$) in each lung compartment during the time interval 0 by t_1 by

$$Y_{\xi}^n = \int_0^{t_1} X_{\xi}^n dt \quad (2.3)$$

where Y_{ξ}^n represents $Y_{\xi}^n(t_1)$.

The cumulated activity for each region of the respiratory system is given by

$$\begin{aligned} U_{NP}^n &= Y_a^n + Y_b^n, \\ U_{TB}^n &= Y_c^n + Y_d^n + Y_k^n + Y_l^n, \\ U_P^n &= Y_e^n + Y_f^n + Y_g^n + Y_h^n, \\ U_L^n &= Y_i^n + Y_j^n, \\ B^n &= Y_a^n \lambda_a^n + Y_c^n \lambda_c^n + Y_e^n \lambda_e^n + Y_i^n \lambda_i^n, \\ G^n &= Y_b^n \lambda_b^n + Y_d^n \lambda_d^n + Y_k^n \lambda_k^n + Y_l^n \lambda_l^n. \end{aligned} \quad (2.4)$$

Here the U_{ξ}^n are the cumulated activity ($\mu\text{Ci-days}$) in the region ξ of the respiratory system, while B^n and G^n represent total activity (μCi) to blood and gastrointestinal tract. In the code, TIMED, the values of Y_{ξ}^n will be computed, but it may be more convenient for the reader to consider the instantaneous state of the system at time t which is represented by the differential equations given in terms of the X 's.

Gastrointestinal Tract

Any exposure by inhalation also leads to some activity entering the gastrointestinal tract (GIT). The dosimetric model for the GIT is essentially due to Eve⁶ so far as the subdivisions of the tract and the transit times through the sections are concerned. Four subdivisions of the tract are defined: the stomach, the small intestine, the upper large intestine, and the lower large intestine. The estimates of dose are considered to be averaged over these sections. Table 2.1 provides data on average masses and times food remains in each of these sections and is quoted from ICRP Publication 23.⁷

Let X_S^n , X_{SI}^n , X_{UL}^n , and X_{LL}^n be the activities present at time t in the contents of the stomach (S), the contents of the small intestine (SI), the contents of the upper large intestine (UL), and the contents of the lower large intestine (LL), respectively, then

$$\begin{aligned} \dot{X}_S^{n+1} &= -X_S^{n+1}(\lambda_S^{n+1} + \lambda_r^{n+1}) + X_b^{n+1}\lambda_b^{n+1} + X_d^{n+1}\lambda_d^{n+1} + X_k^{n+1}\lambda_d^{n+1} \\ &\quad + X_l^{n+1}\lambda_d^{n+1} + X_S^n\lambda_r^{n+1}, \\ \dot{X}_{SI}^{n+1} &= -X_{SI}^{n+1}(\lambda_{SI}^{n+1} + \lambda_r^{n+1} + \lambda_{ab}^{n+1}) + X_S^{n+1}\lambda_S^{n+1} + X_{SI}^n\lambda_r^{n+1}, \quad (2.5) \\ \dot{X}_{UL}^{n+1} &= -X_{UL}^{n+1}(\lambda_{UL}^{n+1} + \lambda_r^{n+1}) + X_{SI}^{n+1}\lambda_{SI}^{n+1} + X_{UL}^n\lambda_r^{n+1}, \\ \dot{X}_{LL}^{n+1} &= -X_{LL}^{n+1}(\lambda_{LL}^{n+1} + \lambda_r^{n+1}) + X_{UL}^{n+1}\lambda_{UL}^{n+1} + X_{LL}^n\lambda_r^{n+1}. \end{aligned}$$

From Table 2.1 $\lambda_S = 24 \text{ days}^{-1}$, $\lambda_{SI} = 6 \text{ days}^{-1}$, $\lambda_{UL} = \frac{24}{13} \text{ days}^{-1}$, and $\lambda_{LL} = 1 \text{ day}^{-1}$. $\lambda_{ab} = \frac{6 \cdot f_1}{1-f_1} \text{ days}^{-1}$ is the fraction of activity present in

Table 2.1. Gastrointestinal Tract Model of Reference Man

Portion of GIT	Mass of Wall (g)	Mass of Contents (g)	Average Time Food Remains (Days)
Stomach	750	250	1/24
Small Intestine	640	400	4/24
Upper Large Intestine	210	220	13/24
Lower Large Intestine	160	135	24/24

the SI which is absorbed to blood per unit time, and f_1 is the fraction of the material ingested which is absorbed to blood.

There are four differential equations corresponding to the GIT for each radionuclide in the chain. For $n=0$, $X_{\xi}^n \equiv 0$ which simplifies the differential equations for the parent. Initial conditions for the GIT following intake by inhalation are $X_{\xi}^n(0) = 0$ for all n and all subscripts ξ . This is easily adapted for oral intake.

The cumulated activity ($\mu\text{Ci-days}$) from time 0 to time t_1 for each section for the GIT is computed by

$$U_{\xi}^n = Y_{\xi}^n = \int_0^{t_1} X_{\xi}^n dt . \quad (2.6)$$

Blood and Other Organs

The activity transferred from the lungs or from GIT to blood will probably deposit in some other organs. For intake of 1 μCi to blood

$$R_j^n(t) = \sum_{s=1}^{q_j^n} a_{js}^n e^{-(\lambda_{js}^n + \lambda_r^n)t} , \quad \begin{array}{l} j = 1, 2, \dots, k, \\ k \equiv \text{the total number} \\ \text{of organs} \end{array} \quad (2.7)$$

represents retention of the j^{th} organ for the n^{th} radionuclide. The coefficients a_{js}^n and the exponential constants λ_{js}^n and λ_r^n are supposedly known. The retention functions $R_j^n(t)$ may be the results of fitting sums of exponentials to retention data on man or on experimental animals. They may also be the results of a compartmental analysis of the problem.

Let X_{js}^n represent retention of the n^{th} radionuclide in the compartment which corresponds to the s^{th} term in the sum for the j^{th} organ. Then

$$\begin{aligned}
\dot{\lambda}_{js}^{n+1} = & -\lambda_{js}^{n+1} (\lambda_{js}^{n+1} + \lambda_r^{n+1}) \\
& + a_{js}^{n+1} (\lambda_a^{n+1} \lambda_a^{n+1} + \lambda_c^{n+1} \lambda_c^{n+1} + \lambda_e^{n+1} \lambda_e^{n+1} + \lambda_i^{n+1} \lambda_i^{n+1} + \lambda_{SI}^{n+1} \lambda_{ab}^{n+1}) \\
& + a_{js}^{n+1} \sum_{l=1}^k \sum_{m=1}^{q_l^n} \lambda_{lm}^n \lambda_r^{n+1} I_{lm}^{n+1} \\
& + \left[\sum_{m=1}^{q_j^n} \lambda_{jm}^n \lambda_r^{n+1} (1 - I_{jm}^{n+1}) \right] a_{js}^n / \sum_{p=1}^{q_j^{n+1}} a_{jp}^{n+1}
\end{aligned} \tag{2.8}$$

where I_{lm}^{n+1} is the fraction of activity of the $n+1$ st daughter that recirculates to blood, and

$$\sum_{j=1}^k \sum_{s=1}^{q_j^n} a_{js}^n = 1 .$$

There are $\sum_{i=1}^r \sum_{j=1}^k q_j^i$ differential equations for other organs for each radionuclide in the chain. For the parent, $\lambda_{js}^0 = 0$ which simplifies the differential equations for the parent. Initial conditions for other organs following intake by inhalation or ingestion are $\lambda_{js}^n(0) = 0$ for all n and subscripts j, s . This is easily adapted for intake by injection.

The cumulated activity from time 0 to time t_1 for some compartment j, s of the body is given by

$$Y_{js}^n = \int_0^{t_1} \lambda_{js}^n dt . \tag{2.9}$$

For a given j these are summed to form

$$U_j^n = \sum_{s=1}^j v_{js}^n \quad (2.10)$$

In many cases equation (2.7) is given for k organs, one of which is termed "other tissues". When this occurs, the following transformation is performed on the values (2.10). Assuming the k^{th} organ is "other tissues", $U_k^n = U_{\text{other}}^n$, "other tissues" is transformed to total body as follows:

$$U_{TB}^n = U_{\text{other}}^n \cdot \frac{M_{TB}}{M_{TB} - \sum_{j=1}^{k-1} M_j} \quad (2.11)$$

where M_{TB} is the mass of the total body and M_j is the mass of organ j .

All other organs $i = 1, \dots, k-1$ are transformed as follows:

$$U_i^n = U_i^n - U_{\text{other}}^n \cdot \frac{M_i}{M_{TB} - \sum_{j=1}^{k-1} M_j} \quad (2.12)$$

III. COMPUTATIONAL CONSIDERATIONS

Suppose we wish to determine the activity (μCi) from inhalation of 1 μCi of a radionuclide which decays to a chain of daughter radionuclides having total length n . The problem may be expressed in the form

$$\dot{\bar{x}} = A\bar{x} \quad (3.1)$$

given the initial condition $\bar{x}(t_0) = \bar{x}(0)$ where the vectors $\dot{\bar{x}}$ and \bar{x} may be partitioned as

$$\dot{\bar{x}} = \begin{bmatrix} \dot{x}^1 \\ \dot{x}^2 \\ \vdots \\ \dot{x}^n \end{bmatrix}, \quad \bar{x} = \begin{bmatrix} x^1 \\ x^2 \\ \vdots \\ x^n \end{bmatrix}, \quad (3.2)$$

and each vector \dot{x}^i and x^i may be written as

$$\dot{x}^i = \begin{bmatrix} \dot{x}_a^i \\ \vdots \\ \dot{x}_L^i \\ \dot{x}_S^i \\ \vdots \\ \dot{x}_{LL}^i \\ \dot{x}_{11}^i \\ \vdots \\ \dot{x}_{kq_k}^i \end{bmatrix} \left\{ \begin{array}{l} \text{Lungs} \\ \\ \text{GIT} \\ \\ \text{Other} \\ \text{Organs} \end{array} \right. , \quad x^i = \begin{bmatrix} x_a^i \\ \vdots \\ x_L^i \\ x_S^i \\ \vdots \\ x_{LL}^i \\ x_{11}^i \\ \vdots \\ x_{kq_k}^i \end{bmatrix} \left\{ \begin{array}{l} \text{Lungs} \\ \\ \text{GIT} \\ \\ \text{Other} \\ \text{Organs} \end{array} \right. \quad (3.3)$$

where k is the number of organs, other than lungs and GIT, for which retention information is available. Thus, the differential equations will be ordered: lungs, GIT, other organs - parent; lungs, GIT, other organs - 1st daughter, etc. Then the matrix A may be partitioned in correspondence with \dot{x} and x as expressed in (3.2) as

$$A = \begin{bmatrix} A_{11} & & & & & & \\ A_{21} & A_{22} & & & & & 0 \\ & A_{32} & A_{33} & & & & \\ & & & \ddots & \ddots & & \\ & & & & \ddots & & \\ 0 & & & & & A_{n,n-1} & A_{n,n} \end{bmatrix} \quad (3.4)$$

where A_{ii} is a lower triangular matrix with negative diagonal elements. $A_{i,i-1}$ is rectangular having the same number of rows as A_{ii} and the same number of columns as $A_{i-1,i-1}$.

In order to better visualize the form of A , consider an example, for inhalation, with a chain of two radionuclides and two organs where

$$R_1^1(t) = a_{11}^1 e^{-(\lambda_{11}^1 + \lambda_r^1)t},$$

$$R_2^1(t) = a_{21}^1 e^{-(\lambda_{21}^1 + \lambda_r^1)t},$$

and

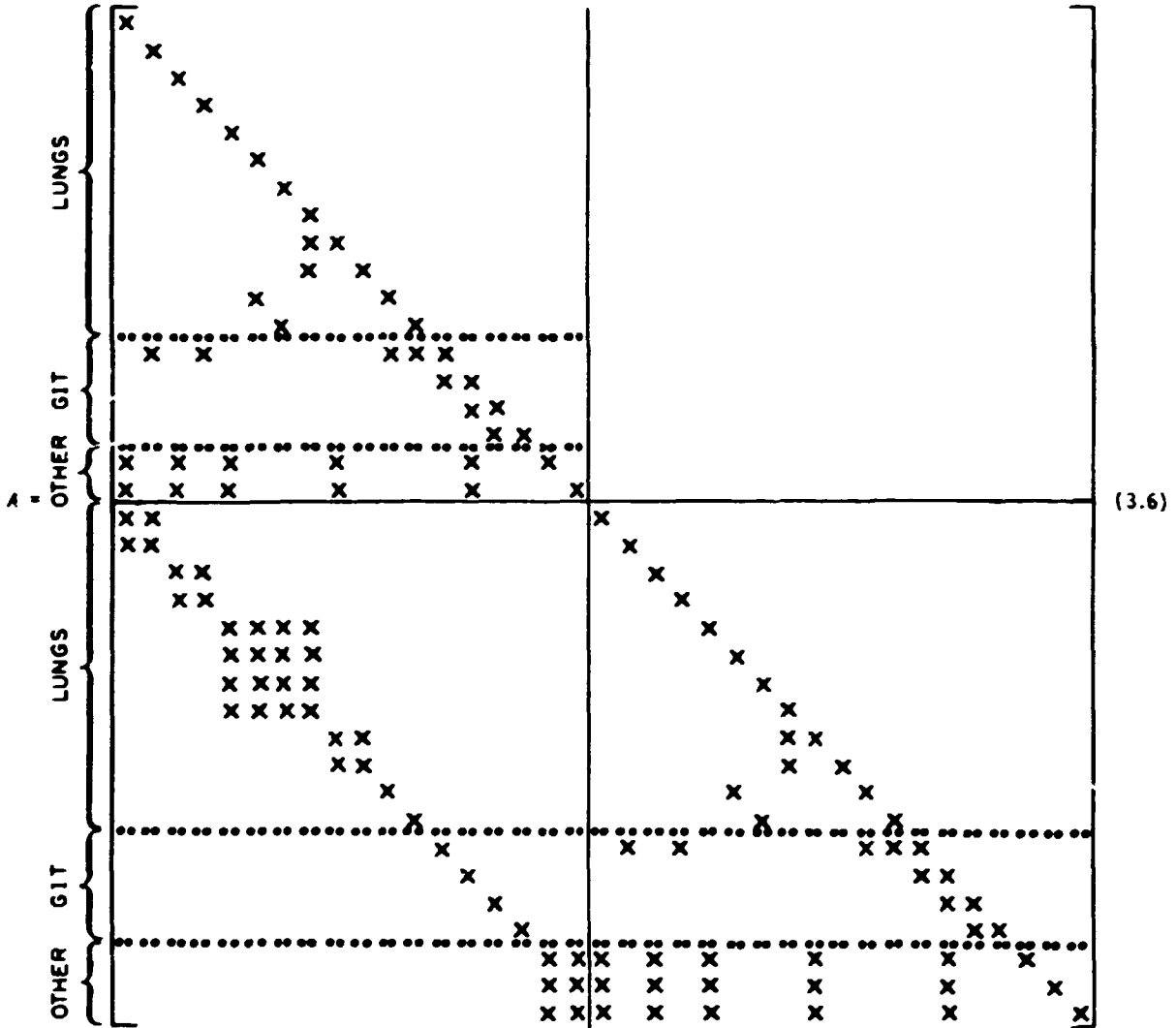
$$R_1^2(t) = a_{11}^2 e^{-(\lambda_{11}^2 + \lambda_r^2)t} + a_{12}^2 e^{-(\lambda_{12}^2 + \lambda_r^2)t},$$

$$R_2^2(t) = a_{21}^2 e^{-(\lambda_{21}^2 + \lambda_r^2)t}.$$

describe retention R_j^i for nuclide i organ j .

There will be 37 (2·12 + 2·4 + 2 + 3) differential equations and matrix A will have the form

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The x's indicate nonzero entries in the matrix; the dotted lines are present to help the reader distinguish visually the coefficients of the terms in the differential equations for the lungs, GIT, and other tissues for both the parent and the daughter; the solid lines indicate where the matrix should be partitioned in order to be of the form

$$A = \begin{bmatrix} A_{11} & 0 \\ A_{21} & A_{22} \end{bmatrix} .$$

Computer Storage of the Coefficient Matrix

It is evident from Eq. (3.6) that the coefficient matrix A is a large, sparse (i.e., few nonzero entries) matrix. For computational purposes the matrix is stored in the computer as three vectors. The first vector contains the nonzero elements of A stored by rows. The second and third vectors hold the row and column index, respectively, for each of the nonzero elements of A . The method of storage is required due to the fact that even for a short chain of radionuclides the coefficient matrix becomes too large to store reasonably the entire matrix in the computer.

Other Modes of Intake

Ingestion or injection of 1 μCi of a radionuclide may be achieved by a slight modification of the system of differential equations. For ingestion, remove from the system those equations for the lungs and modify the initial conditions so that 1 μCi is deposited in the stomach.

$$x_S^1(0) = 1.0 \quad , \quad (3.7)$$

$$x_{SI}^1(0) = x_{UL}^1(0) = x_{LL}^1(0) = 0.0 \quad .$$

For injection, remove from the system those equations describing lung and GIT retention, and alter the initial condition vector so that

$$x_{js}^1(0) = a_{js}^1 / \sum_{i=1}^k \sum_{t=1}^{q_i^1} a_{it}^1 \quad (3.8)$$

where k is the number of organs and q_i^1 is the number of terms in the retention function corresponding to the i^{th} organ. Thus for ingestion and injection (3.3) reduces to (3.9) and (3.10), respectively,

$$\text{for ingestion - } \dot{x}^i = \left[\begin{array}{c} \dot{x}_S^i \\ \vdots \\ \dot{x}_{LL}^i \\ \dot{x}_{11}^i \\ \vdots \\ \dot{x}_{kq_k}^i \end{array} \right] \left. \begin{array}{l} \text{GIT} \\ \\ \text{Other} \\ \text{Organs} \end{array} \right\} , \quad x^i = \left[\begin{array}{c} x_S^i \\ \vdots \\ x_{LL}^i \\ x^i \\ \vdots \\ x_{kq_k}^i \end{array} \right] \left. \begin{array}{l} \text{GIT} \\ \\ \text{Other} \\ \text{Organs} \end{array} \right\} \quad (3.9)$$

$$\text{for injection - } \dot{x}^i = \left[\begin{array}{c} \dot{x}_{11}^i \\ \vdots \\ \dot{x}_{kq_k}^i \end{array} \right] \left. \begin{array}{l} \text{Other} \\ \text{Organs} \end{array} \right\} , \quad x^i = \left[\begin{array}{c} x_{11}^i \\ \vdots \\ x_{kq_k}^i \end{array} \right] \left. \begin{array}{l} \text{Other} \\ \text{Organs} \end{array} \right\} \quad (3.10)$$

For the example shown in Eq. (3.6) the matrix A would be reduced to

$$A = \begin{array}{c} \left. \begin{array}{l} \text{OTHER GIT} \\ \text{OTHER GIT} \\ \dots \\ \text{OTHER GIT} \\ \text{OTHER GIT} \end{array} \right\} \begin{array}{|c|c|} \hline \begin{array}{c} \times \times \\ \times \times \\ \dots \times \times \dots \\ \times \times \\ \times \end{array} & \\ \hline \begin{array}{c} \times \\ \times \\ \dots \times \dots \\ \dots \times \times \dots \\ \times \times \\ \times \times \\ \times \times \end{array} & \begin{array}{c} \times \\ \times \times \\ \times \times \\ \dots \times \times \dots \\ \times \times \\ \times \\ \times \end{array} \\ \hline \end{array} \quad (3.11)$$

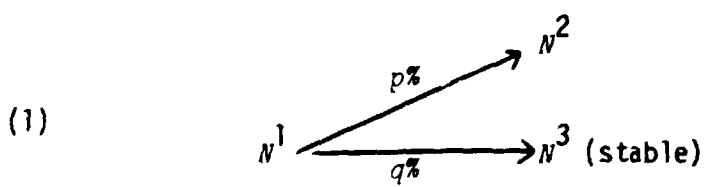
for ingestion, and

$$A = \begin{array}{c} \left. \begin{array}{l} \text{OTHER} \\ \text{OTHER} \\ \dots \\ \text{OTHER} \\ \text{OTHER} \end{array} \right\} \begin{array}{|c|c|} \hline \begin{array}{c} \times \\ \times \end{array} & \\ \hline \begin{array}{c} \times \times \times \\ \times \times \times \\ \times \times \end{array} & \begin{array}{c} \times \\ \times \\ \times \end{array} \\ \hline \end{array} \quad (3.12)$$

for injection.

Branching in the Chain of Radionuclides

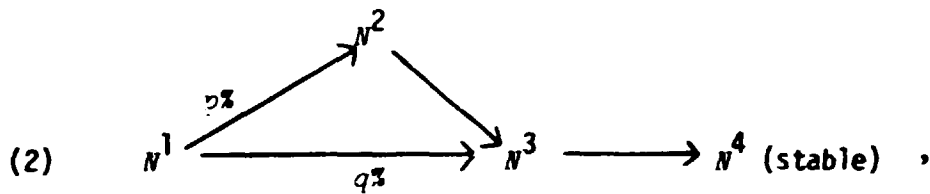
When the decay chain involves branching, the coefficient matrix takes a different form than that shown in Eq.(3.4). The authors have considered three forms of branching. By numbering the nuclides as they will be ordered in Eq. (3.2), we have



Then the partitioned matrix A takes the form

$$A = \begin{bmatrix} A_{11} & 0 \\ \frac{p}{100} \cdot A_{21} & A_{22} \end{bmatrix} \quad (3.13)$$

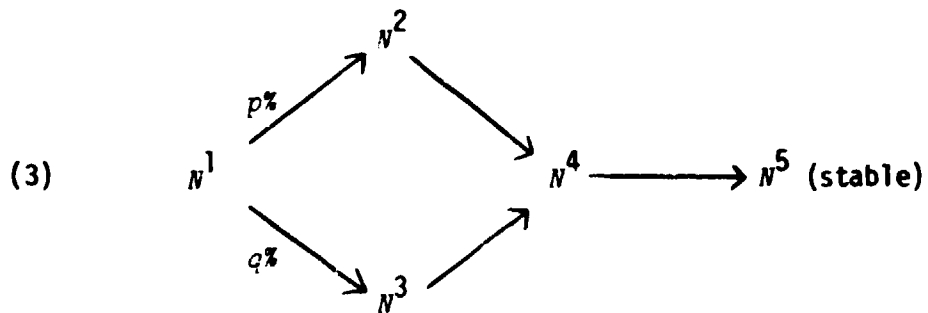
where $\frac{p}{100} \cdot A_{21}$ indicates that every element in the matrix A_{21} is multiplied by $\frac{p}{100}$. If a chain has the form



then

$$A = \begin{bmatrix} A_{11} & 0 & 0 \\ \frac{p}{100} \cdot A_{21} & A_{22} & 0 \\ \frac{q}{100} \cdot A_{31} & A_{32} & A_{33} \end{bmatrix}. \quad (3.14)$$

The chain



has the corresponding coefficient matrix:

$$A = \begin{bmatrix} A_{11} & 0 & 0 & 0 \\ \frac{p}{100} \cdot A_{21} & A_{22} & 0 & 0 \\ \frac{q}{100} \cdot A_{31} & 0 & A_{33} & 0 \\ 0 & A_{42} & A_{43} & A_{44} \end{bmatrix} . \quad (3.15)$$

Certainly these branching schemes do not stand alone or occur only at the beginning of a chain. Thus, the coefficient matrix A will, in general, be some combination of (3.4) and (3.13), (3.14), or (3.15). These examples have sufficed to the present, but it is possible other forms may occur. However, the general pattern seems clear from these examples.

Calculation of Cumulated Activity

It should be noted that, in general, we do not wish to obtain the activity (μCi) as the final result but cumulated activity ($\mu\text{Ci-days}$) which represents the time integral of the solution to (3.1). This is done in a very straightforward manner.

Given $\dot{x}(t) = AX(t)$, Eq. (3.1), with initial condition $x(0) = x_0$ and integrating both sides we have

$$\int_0^{t_1} \dot{x}(t) dt = \int_0^{t_1} AX(t) dt ,$$

$$x(t_1) - x_0 = A \int_0^{t_1} x(t) dt ,$$

$$x(t_1) = A \int_0^{t_1} x(t) dt + x_0 .$$

By letting $Y(t_1) = \int_0^{t_1} X(t) dt$ where $Y(0) = 0$.

$$\dot{Y}(t_1) = \left[\int_0^{t_1} X(t) dt \right]' = X(t_1) .$$

Thus we may reformulate (3.1) as

$$\dot{Y} = AY + X_0 \quad \text{where } Y(0) = 0 , \quad (3.16)$$

and (3.16) will be solved for Y to determine the cumulated activity ($\mu\text{Ci-days}$) in that organ from 0 to t_1 . It is Eq. (3.16), not Eq. (3.1), to which TIMED calculates the solution.

Solution of the Differential Equations

The solution of (3.16) is calculated by using a FORTRAN subroutine package written by A. C. Hindmarsh,⁸ based on a program written by C. W. Gear,⁹ for the solution of the initial value problem for system of ordinary differential equations (ODE's). In general, such a system has the form

$$\dot{y} = f(y, t) \quad (3.17)$$

or more specifically

$$\frac{dy_i(t)}{dt} = f_i(y_1(t), \dots, y_N(t), t) \quad (3.18)$$

where y , \dot{y} , and f are vectors of length $N \geq 1$. Given an initial value of the vector

$$y(t_0) = y_0,$$

and a subroutine for the calculation of f , the GEAR package can be used to compute a numerical solution to (3.18) at values of the independent variable t in some interval (t_0, T) , as desired by the user.

The basic methods used for the solution are of implicit linear multistep type. There are classes of such methods available to the user. The first is the implicit Adams methods (up to order 12), and the second is the backward differentiation formula (BDF) methods (up to order 5), also called Gear's stiff methods.

A prime feature of GEAR is the ability to solve stiff ODE problems. Roughly speaking, an ODE system is called stiff if it involves both very rapidly changing terms and very slowly changing terms, all of a decaying nature. More precisely, we consider the eigenvalues v_i of the $N \times N$ Jacobian matrix

$$J = \frac{\partial f}{\partial y} = \left(\frac{\partial f_i}{\partial y_j} \right)_{i,j=1}^N \quad (3.19)$$

and suppose that the v_i all have negative real parts. The "time constants" of the problem are then $\tau_i = 1/|\operatorname{Re}(v_i)|$, and the decaying nature (locally) of the solution is given by the exponentials e^{-t/τ_i} .

If the N time constants τ_i are widely spread, and those terms with the smaller τ_i have already decayed to an insignificant level, then the system is stiff.

upon considering the properties of Eq. (3.16), it can be seen that the Jacobian of the matrix A is itself (that is, $J(A) = A$), and, A being a nonsingular, lower triangular matrix, the eigenvalues of A are the diagonal elements of the matrix, which are $-(\lambda_{\xi}^j + \lambda_r^j)$. Thus the time constants τ_i are

$$\tau_i = \frac{1}{\lambda_{\xi}^j + \lambda_r^j}, \quad i = 1, \dots, N$$

where ξ indicates the compartment and j indicates the member of the chain. The τ_i are of greatly differing orders of magnitude, due to the differing values of the radiological decay constants, λ_r^j , for each nuclide j , and also the differing biological decay constants, λ_{ξ}^j , for various compartments ξ . This implies the system is stiff.

The difficulty with stiff problems is that most conventional methods for solving ODE's require incremental values of t commensurate with $\min \tau_i$, while the size $|T-t_0|$ of the problem range is commensurate with $\max \tau_i$. As a result, the problem cannot be run to completion in a reasonable number of steps. With Gear's methods, however, the time increment h is restricted to small values, by the requirement of accuracy, only where the solution is relatively active. By definition, the problem is not stiff in such regions, and accuracy is achieved at minimum computational cost by allowing both h and the order of the method to vary. Then in regions of stiffness, where the solution is inactive, Gear's methods

have the property of "stiff stability", which assures that h is no longer restricted by small time constants, unless or until the corresponding, rapidly decaying terms become active again.

The GEAR package also contains, as an option, a method well suited for nonstiff problems, namely the implicit Adams methods with fixpoint corrector iteration, also called the Adams-Bashforth-Moulton methods. Both the stiff and nonstiff methods are implemented in a manner which allows both the step size and the order to vary in a dynamic way throughout the problem. This variability is now widely recognized as highly desirable for efficiency in using linear multistep methods.

The methods used in the GEAR package are documented in considerable detail elsewhere.^{9,10,11} Hence, only a brief summary of them will be given here.

The basic methods involved are linear multipoint methods of the form

$$y_n = \sum_{j=1}^{K_1} \alpha_j y_{n-j} + h \sum_{j=0}^{K_2} \beta_j \dot{y}_{n-j}, \quad (3.20)$$

where y_k is an approximation to $y(t_k)$, $\dot{y}_k = f(y_k, t_k)$ is an approximation to $\dot{y}(t_k)$, and h is a constant step size: $h = t_{k+1} - t_k$. In the case of the Adams method of order q , we have $K_1 = 1$ and $K_2 = q - 1$. In the case of the backward differentiation formula (BDF) of order q , we have $K_1 = q$ and $K_2 = 0$. The BDF's are so called because, on dividing through by $h\beta_0$, they can be regarded as approximation formulas for \dot{y}_n in terms of $y_n, y_{n-1}, \dots, y_{n-q}$. In either case, the α_j and β_j are constants

associated with the method and $\beta_0 > 0$. The latter means that Eq. (3.20) is an implicit equation for y_n and is, in general, a nonlinear algebraic system that must be solved at every step. The fact that the order of a given method is q means that, if Eq. (3.20) is solved for y_n with all past values being exact, then y_n will differ from the correct solution of the ODE by a local truncation error that is of order h^{q+1} , commonly expressed as $O(h^{q+1})$, for small h .

If Eq. (3.20) is written in the form

$$g(y_n) \equiv y_n - \beta_0 f(y_n, t_n) - \sum_1^{K_1} \alpha_j y_{n-j} - h \sum_1^{K_2} \beta_j \dot{y}_{n-j} = 0, \quad (3.21)$$

then the nonlinear system $g(y_n) = 0$ can be solved, for example, by Newton's method:

$$y_{n(m+1)} = y_{n(m)} - P_{n(m)}^{-1} g(y_{n(m)}), \quad (3.22)$$

$$P_{n(m)} = \left. \frac{\partial g}{\partial y} \right|_{y_{n(m)}} = I - h \beta_0 \left. \frac{\partial f}{\partial y} \right|_{y_{n(m)}}.$$

The manner in which past values are saved is rather unusual by comparison with most other ODE programs and was invented by A. Nordsiech.¹² While the conventional choice would be to store an array of $L = K_1 + K_2 + 1 = q + 1$ current and past values of y_k and \dot{y}_k , Nordsiech's history is a linear transform of this one and has the form

$$\underline{z}_n = (y_n, h \dot{y}_n, h^2 \ddot{y}_n / 2, \dots, h^q y_n^{(q)} / q!). \quad (3.23)$$

This is an $N \times L$ array of approximate "scaled derivatives" of y at t_n up to order q and is the Y array of the program.

With this choice of history array, the prediction of \underline{z}_n (and in fact all of \underline{z}_n) from \underline{z}_{n-1} becomes quite simple. It is given by

$$\underline{z}_n(0) = \underline{z}_{n-1}A, \quad A = \left(\binom{i}{j} \right)_{i,j=0}^q, \quad \text{that is } \begin{cases} a_{ij} = \frac{i!}{j!(i-j)!} & i \geq j \\ a_{ij} = 0 & i < j \end{cases} \quad (3.24)$$

A is the $L \times L$ Pascal triangle matrix. As Gear observed, the multiplication $\underline{z}A$, for a row vector $\underline{z} = (z_i)_{i=0}^q$ can be accomplished with additions only, as follows:

$$\underline{z}_{j-1} \leftarrow \underline{z}_{j-1} + \underline{z}_j \quad (3.25)$$

where $j = q, q-1, \dots, i-1$ and $i = 0, 1, \dots, q-1$.

The complete algorithm for the prediction and corrections at step n can then be written as follows:¹⁰

$$\begin{aligned} \underline{z}_n(0) &= \underline{z}_{n-1}A, \\ \left. \begin{aligned} y_{n(m+1)} &= y_{n(m)} + \ell_0 P_n^{-1} F_m, \\ h\dot{y}_{n(m+1)} &= h\dot{y}_{n(m)} + P_n^{-1} F_m, \\ F_{(m)} &\equiv hf(y_{n(m)}, t_n) - h\dot{y}_{n(m)}, \end{aligned} \right\} (m=0, 1, \dots, M-1) \quad (3.26) \end{aligned}$$

$$y_n = y_{n(M)} = y_{n(0)} + \ell_0 E_n,$$

$$E_n = \sum_{m=0}^{M-1} P_n^{-1} F_m,$$

$$\underline{z}_n = \underline{z}_n(0) + E_n \ell.$$

Here M is the number of iterations performed, and $\underline{\lambda} = (\lambda_i)_0^q$ is a row vector of constants determined by the basic method, Eq. (3.20), and satisfying $\lambda_0 = \beta_0$ and $\lambda_1 = 1$. The vector E_n is saved, as it is proportional to the estimated truncation error committed on the step.¹¹

Following a step of size h at order q , the GEAR package, at intervals of $q + 2$ steps, is programmed to choose a larger step size by estimating the local truncation errors at orders $q-1$, q , and $q+1$. The largest value h' of the three step sizes obtained is then chosen, and the order reset accordingly. Also the Nordsiech array must be rescaled by powers of h'/h . The data used to take the subsequent steps of size h' is in effect obtained by interpolating with the data at a spacing of h .

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IV. PROGRAM DESCRIPTION

The program TIMED consists of sixteen routines that are programmed in a fairly well-structured manner and are heavily augmented with program comment cards. Hence, to anyone who is familiar with the methods used in this report and in Refs. 8, 9, 10, and 11, the details are self-explanatory. However, to supplement this material, we include the diagram of program structure and a description of each routine.

Fig. 4.1 shows the overall structure of the TIMED program. Arrows indicate the calling sequence of the routines. Routines DRIVES, STIFF, YOUT, YOUTD, YNA, and TESBC have been taken from the GEAR package⁸ and have been specialized for use in TIMED by D. E. Arnarius at Oak Ridge National Laboratory.¹³ MAIN, YPAY, and AXEB have been written by him for use on this particular problem with our computer, the IBM System/360. The remainder of the routines have been written by one author (S. B. Watson) to implement the model described in Section II.

Except for AXEB and YPAY, all routines are written in the IBM System/360 and System/370 FORTRAN IV language. AXEB and YPAY are IBM System/360 Assembler language routines. A listing of TIMED routines is given in Appendix A.

The MAIN routine functions as follows:

1. Read the values N_0 , T_{LAST} , H_1 , EPS .

N_0 - the number of first order differential equations.

T_{LAST} - the final value of T , i.e. integration proceeds from $T = T_0$ to $T = T_{LAST}$.

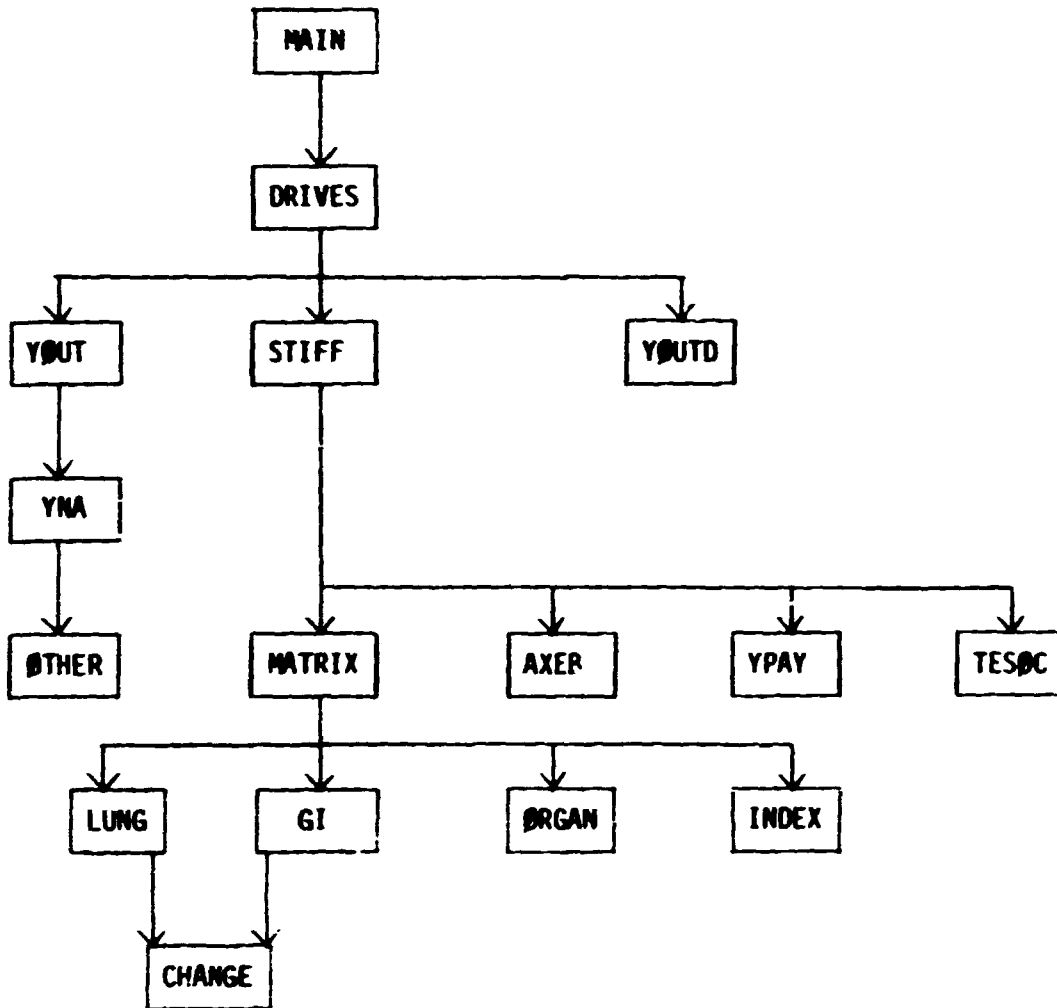


Fig. 4.1. Overall Structure of TIMED.

H1 - initial step size for the integration procedure.
 EPS - the local error tolerance parameter. The estimated errors δ_i in y_i are compared to $YMAX(i)$, which is roughly the largest value of $|y_i|$ seen so far, for error control purposes this is done by comparing the root-mean-square (RMS) norm (i.e. Euclidean norm divided by \sqrt{N} of the vector $[\delta_i/YMAX(i)]_{i=1}^N$) to EPS. This ratio is kept less than EPS.

2. Read the number of print intervals, NPRT, and the arrays defining the print times, TPRL, DTPR, and NARPRT by

```
DO 10 I = 1, NPRT
```

```
10 TPRINT = TPRL(I) + Q*DTPR(I)
```

where $Q = 0, \dots, NARPRT(I)$.

3. Check to see if $NPRT \leq 10$.
4. Check to see if $TPRL(NPRT) + NARPRT(NPRT)*DTPR(NPRT) = TLAST$.
5. Check to see that consecutive time intervals do not overlap.
6. Set TPRINT to TPRL(1,).
7. Check to see that $13*N0 \leq 2400$ for Adams procedure (i.e. $N0 \leq 125$) or $6*N0 \leq 2400$ for stiff procedure (i.e. $N0 \leq 400$).
8. Call DRIVES.
9. STOP.

The routine DRIVES is an interface between the user and the rest of the GEAR package. It oversees the integration of the ODE over the interval between two of the user's output points. DRIVES proceeds as follows:

1. Test input parameters for correctness and initialize variables.
2. Call STIFF.
3. KFLAG set in STIFF and returned to DRIVES as one of the following values:
 - 0 problem was completed successfully.
 - 1 the integration was halted after failing to pass the error test even after reducing H by a factor of 10^{10} from its initial value.
 - 2 after some initial success, the integration was halted when a reduction in H by a factor of more than 10^4 was indicated in order to pass the error test.
 - 3 the integration was halted after failing to achieve corrector convergence even after reducing H by a factor of 10^{10} from its initial value.
 - 4 immediate halt because of illegal values of input parameter.
 - 5 H is such that $T + H = T$.
4. If KFLAG = 0 or -4, call YOUT for normal output of Y at time TPRINT; go to 2.
5. If KFLAG = -1, print message; try 10 more reductions of H; call YOUTD if continues to fail; return.
6. If KFLAG = -2, -3 -5, print error message; call YOUTD; return.

YOUT computes interpolated values of the dependent variable Y , and calls YNA to print these values. YNA prints the solution vector Y , Eq. (3.16), and also computes Eqs. (2.4), (2.6), and (2.10) and prints these quantities. YNA calls the routine OTHER to perform the "other tissues" transformation described in (2.11) and (2.12), should it be required. YOUTD is a dummy routine which could be expanded by the user to print diagnostic information.

STIFF performs one step of the integration of an initial value problem for a system of ordinary differential equations. Fig. 4.2 shows the general flow structure of the routine.

Recalling the unusual structure of the coefficient matrix, A , in (3.1) as discussed in Section III, A is stored as a vector which is limited in length by STIFF to a maximum of 6000 elements. This means that the user is limited to 6000 nonzero elements in the coefficient matrix. A check is made to see that sufficient storage is available for A . STIFF also counts the number of nonzero elements in each row of A and writes an error message if this number exceeds 52. It should be noted that some of the methods for storing elements in certain arrays are both compiler- and machine-dependent. This may cause problems on machines other than IBM System/360 or System/370.

AXEB is an IBM 360 Assembler language routine called by STIFF to solve the matrix equation $AX = b$ where A is an N by N lower triangular matrix, and A is stored as a vector consisting of the nonzero entries in A stored sequentially by rows. It is assumed that the maximum number of nonzero entries in a row of A is less than or equal to 52.

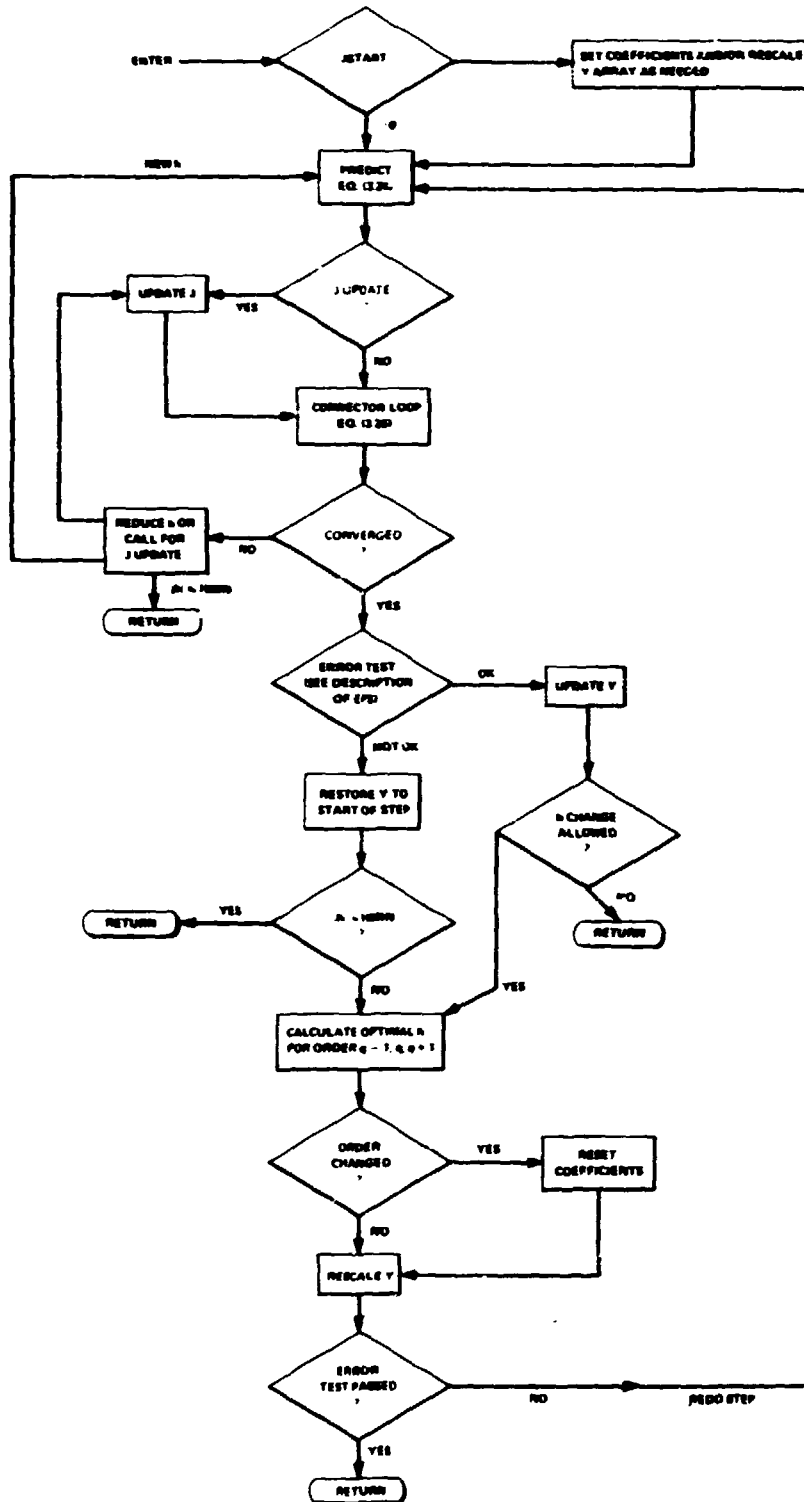


Fig. 4.2. Macroscopic Flow Chart of STIFF (Taken from Ref. 8).

YFAY is also an Assembler language routine. It is called by STIFF to form $\dot{Y} = AY + X_0$. The matrix A is stored as in AXEB, and this routine assumes a maximum of 52 nonzero entries.

AXEB and YPAY were written so as to most efficiently make use of the properties of the matrix A and of the computer storage.

TESOC is called by STIFF and sets the coefficients that are used there, both for the basic integration step and for error control. TESOC also sets MAXDER, the maximum order of the method available. Currently it is 12 for Adam's method and 5 for the GEAR method.

MATRIX directs the formation of the coefficient matrix, A , and may be outlined as follows:

1. Read descriptive information about the chain of radionuclides.

NISØ - number of radionuclides in the chain

D3, D4, D5 - regional deposition values⁴

NØRG - number of organs (excluding lung and GIT)

IØPT - mode of intake (0 - inhalation, 1 - ingestion,
2 - injection)

2. Call LUNG to assign regional deposition for each pathway a,...,h (see Fig. 2.1 for explanation of lung pathways).

3. Read for each radionuclide

TAG - atomic number and symbol

TR - half life

ICLASS - inhalation class

ILAB - (0 - absorption in SI only, 1 - absorption in
other regions of GIT)

4. Set LAMR.

$$\text{LAMR} = \ln 2 / \text{TR}$$

5. Call LUNG1 to set LAMX and FRAC according to inhalation class.

$$\text{LAMX} = \lambda_{\xi} = \ln 2 / T_{\xi}, \quad T_{\xi} \text{ is the removal half time described in Fig. 2.1}$$

$$\text{FRAC} = F_{\xi} \text{ is the compartmental fraction from Fig. 2.1}$$

6. Initialize row and column indices.

7. For each nuclide:

(a) If intake by inhalation, call LUNG2 to set up portion of matrix corresponding to lung differential equations.

(b) If intake is by inhalation or ingestion, call routine GI to set up portion of matrix corresponding to GIT differential equations.

(c) Call ORGAN to set up coefficient matrix for differential equations corresponding to organs other than lungs and GIT for which we have retention models.

8. Print coefficient matrix.

9. Print nonzero elements of the coefficient matrix and the row and column in which each one appears. Recall that, due to the sparse nature of the coefficient matrix, it is stored as three vectors - A, RINDEX, and CINDEX. A(I) is a nonzero element of the coefficient matrix, RINDEX(I) contains the row of A(I), and CINDEX(I) is the column index of A(I). The coefficient matrix is filled from left to right, one row at

a time, proceeding from top to bottom of the matrix, so that the vector A is formed correctly. Note that the components of the vector RINDEX form an increasing sequence, i.e., $RINDEX(I) < RINDEX(I+1)$ while the components of CINDEX do not.

Routine LUNG has three entry points which are called from MATRIX and function as follows:

Entry LUNG - assigns the regional deposition values for each pathway (these are used in setting the initial values for inhalation, see Eq. (2.2)).

Entry LUNGI - determines inhalation class and sets λ_{ξ} and F_{ξ} from Fig. 2.1 which appear in Eq. (2.1). LUNGI also sets the initial values, Eq. (2.2).

Entry LUNG2 - forms the coefficient matrix for Eq. (2.1).

Routine GI is called from MATRIX and determines the entries of the coefficient matrix corresponding to the compartments of the GIT, Eq. (2.5). If intake is by ingestion, initial conditions are set as in Eq. (3.7).

ORGAN functions as follows:

1. For each organ, read organ number (ISORS), the organ name (ORGNM), and the number of compartments (ICOM); i.e., the number of terms in the retention function (see Eq. (2.7)), and the retention function.
2. Form the entries of the coefficient matrix corresponding to Eq. (2.8) for each compartment.

3. If intake is by injection, set the initial values according to Eq. (3.8).

INDEX is called by MATRIX and determines the first column of the matrix entries for each daughter. (Recall the way the matrix entries are filled – from left to right and top to bottom.) This is especially important when branching is considered. Whereas daughter entries begin in a regularly indented fashion for nonbranching radionuclides, Eq. (3.4), the daughter entries have a somewhat irregular pattern in branching chains, see Eqs. (3.13), (3.14), and (3.15).

CHANGE is called by LUNG and GI and determines the initial column for each row of the coefficient matrix for the equations corresponding to N^3 , Eq. (3.14), and the equations corresponding to N^4 , Eq. (3.15).

V. USER INFORMATION

TIMED is written in the IBM System/360 and System/370 FORTRAN IV language with the exception of two routines (AXEB and YPAY which are discussed in Section IV) that are written in IBM System/360 Assembler language. The FORTRAN routines have been compiled using the FORTRAN H-level compiler; the Assembler routines have been compiled with the Assembler F-level compiler. TIMED has been executed on both the IBM 360/91 and 360/75 at ORNL. The program requires about 310 K (1K = 1024 bytes; 1 byte = 8 bits) of core storage for execution. The FORTRAN routines require about 7.5 seconds to compile; the Assembler routines require 0.7 seconds for compilation. Execution time depends upon many factors: the number of differential equations, the number of nonzero entries in the coefficient matrix, the length of time period (t_0, t_{final}) for integration, etc. The execution time for the second example in Appendix B, a complicated chain of four radionuclides involving 116 differential equations, is about 5.8 seconds.

The remainder of this section describes the content of the card input which must be prepared in order to execute the program and the output produced by the program. Three examples of sample input and the corresponding output are presented in Appendix B.

Input

The card input data are to be prepared in the order shown in Table 5.1. Referring to Table 5.1, "Subroutine" indicates the subroutine in

Table 5.1. Card Input to TIMED

Subroutine	Card Number	Columns	Remarks, Variables, Etc.
<u>Information Concerning the Differential Equations</u>			
MAIN	1		FORMAT (15, 7E10.2)
		1-5	NO - order of lower triangular matrix A , that is, the number of differential equations.
		6-15	TLAST - integrate $Y'(t) = AY(t) + X_0$ from $t = 0$ to $t =$ TLAST. TLAST given in units of days.
		16-25	H1 - initial step size for the integration procedure. If left blank, then default is $H1 = 10^{-7}$.
		26-35	EPS - controls the local truncation error during the integration procedure. If left blank, then default is $EPS = 10^{-6}$.
	2		FORMAT (15)
		1-5	NPRT - number of print intervals, $NPRT \leq 10$.
	2+I, I=1,...,NPRT		FORMAT (2E10.2, 15)
		1-10	TPRL(I) - first print time in this print interval.
		11-20	DTPR(I) - print time step for this interval.
		21-25	NARPR(I) - number of time steps to use in the current interval.

Table 5.1. (cont'd)

Subroutine	Card Number	Columns	Remarks, Variables, Etc.
TPRL, DTPR, and NARPRT are defined by:			
$DQ \text{ TO } I = 1, \text{ NPRT}$ $\text{TO TPRINT} = \text{TPRL}(I) + Q * \text{DTPR}(I) \quad Q = 0, \dots, \text{NARPRT}(I)$			
The values assumed by TPRINT must form an increasing sequence.			
<u>General Information for Chain of Radionuclides</u>			
MATRIX	1		FORMAT (I10, 3F10.3, 3I10)
		1-10	NISØ - number of isotopes in the chain, NISØ < 10.
		11-20	D3 - fraction of intake deposited in the NP region.
		21-30	D4 - fraction of intake deposited in the TB region.
		31-40	D5 - fraction of intake deposited in the P region.
		41-50	NØRG - number of organs (excluding lungs and GIT), NØRG < 10.
		51-60	IØPT - mode of intake. 0 - inhalation 1 - ingestion 2 - injection
		61-70	NØRAL - number of ingestion cases to be run (used only with ICRP work). Leave blank if IØPT ≠ 1.
	1+1, I=1, ..., NISØ		FORMAT (A8, 2X, F10.0, 8X, A2, 2F10.0, 2I2, 2X, I2)
	1-8	TAG(I) - Atomic symbol and atomic number of ITH member of the chain.	

Table 5.1. (cont'd)

Subroutine	Card Number	Columns	Remarks, Variables, Etc.
		11-20	TR(I) - half life of ITH member of chain in days.
		29-30	ICLASS(I) - inhalation class of ITH member of chain. Should be one of the characters D, W, or Y right justified in the field. Leave blank if IOPT \neq 0.
		31-40	LAMB(I,2) - absorption coefficient in the small intestine (SI) $\lambda_{ab} = \frac{6 \cdot 10^{-4}}{1 - f_1}$
		41-50	P(I) - fraction of decay to radionuclide I.
		51-52	NBRNCH(I) - type of branching (1, 2, 3).
		53-54	IBRNCH(I) - position in the branching scheme (1, 2, 3).

Three different types of branching cases have been programmed, and the type of branching and position in the branching scheme are defined by the ordered pair (NBRNCH(I), IBRNCH(I)). The ordered pairs are described in the diagrams below.

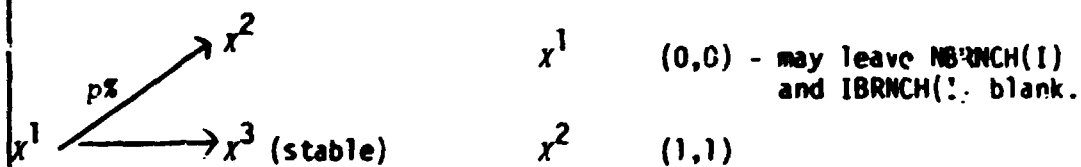


Table 5.1. (cont'd)

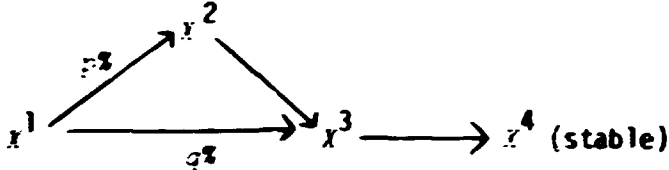
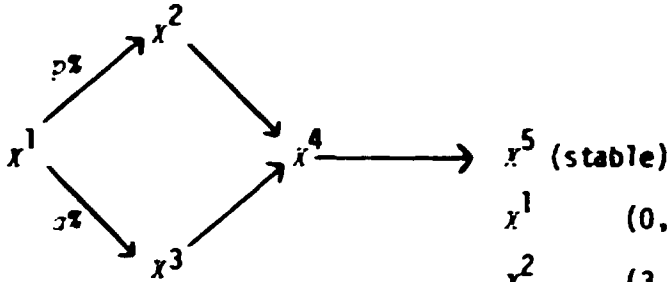
Subroutine	Card Number	Columns	Remarks, Variables, Etc.
 <pre> graph LR x1 -- p% --> x2 x1 -- q% --> x3 x2 --> x3 x3 --> x4["x4 (stable)"] </pre>			
			x^1 (0,0) x^2 (2,1) x^3 (2,2)
 <pre> graph LR x1 -- p% --> x2 x1 -- q% --> x3 x2 --> x4 x3 --> x4 x4 --> x5["x5 (stable)"] </pre>			x^1 (0,0) x^2 (3,1) x^3 (3,2) x^4 (3,3)
		57-58	ILAB - switch to indicate if there is absorption in some section of the GIT tract other than the small intestine (SI) 0 - absorption only in small intestine. 1 - absorption in stomach(S), upper large intestine (UL), or lower large intestine (LL) in addition to small intestine absorption.

Table 5.1. (cont'd)

Subroutine	Card Number	Columns	Remarks, Variables, Etc.
	2-1+i I=1,...,NIS		FORMAT (3F10.0)
Card required only if ILAB = 1.			
		1-10	LAMB(I,1) - adsorption coefficient for S.
		11-20	LAMB(I,3) - adsorption coefficient for UL.
		21-30	LAMB(I,4) - adsorption coefficient for LL.
<u>Description of Retention</u>			
This set of cards describes retention in those organs other than the lungs and GIT. These cards must be included for each radionuclide in the chain.			
ORGAN	1		FORMAT (10I2)
This card should be omitted for the first radionuclide in the chain.			
		1-2	IS(1) - fraction of daughter radionuclide which returns to blood for the 1st organ. (1-IS(1)) remains in the organ.
		3-4	IS(2) - same fraction for 2nd organ.
		⋮	⋮
Only NORG numbers should be placed on the card.			

Table 5.1. (cont'd)

Subroutine	Card Number	Columns	Remarks, Variables, Etc.
Include cards 2 and 3 for each organ.			
	2	1-2	FORMAT (12, A8, 12) ISORS(I) - index of organ I taken from the following list. ISORS(I) is used for indexing purposes when calculating dose.
1. bladder contents		13. ovaries	
2. stomach contents		14. pancreas	
3. SI + content		15. cortical bone (trabecular bone)	
4. UL		16. cancellous bone	
5. LL		17. red marrow	
6. kidneys		18. yellow marrow	
7. liver		19. skin	
8. salivary gland		20. spleen	
9. TB region		21. testes	
10. P region		22. thyroid	
11. lymph		23. total body (used for other tissues, also)	
12. muscle		25. adrenals	
(24. cloud should not be used as a source organ in this code.)			
		3-10	ORGN(I) - name of organ I (at most 8 characters).
		11-12	ICOM(K,I) - number of compartments for organ I, radio-nuclide K (number of terms in the retention function), $ICOM(K,I) \leq 10$.
	3		FORMAT (8F10.0)
This card describes the retention function and contains the coefficients of the exponential, $AS(J,I), JJ=1, \dots, ICOM(K,I)$, and the exponential constants, $TBS(JJ,I), JJ=1, \dots, ICOM(K,I)$, respectively. Obviously, for more than four compartments, i.e., terms in the retention function, another card is needed.)			

Table 5.1. (cont'd)

Subroutine	Card Number	Columns	Remarks, Variables, Etc.
<p>In reference to cards 2 and 3, cards describing total body or "other tissues" should be placed last.</p>			
<p>For each nuclide the organs and associated information must appear in the same order.</p>			
<p><u>Other Tissues Transformation</u></p>			
<p>If one of the source organs is "other tissues", the cumulated activity for these source organs must be transformed as described in Section II. To invoke the procedure, a card must be included for each print interval of the form:</p>			
<p>OTHER</p>	<p>1</p>	<p>1-2 3-4 5-6 ⋮</p>	<p>FORMAT (10I2) NØTØP - nonzero integer IØTHER(1) - ISØRS(1) IØTHER(2) - ISØRS(2) ⋮</p>
<p>Do not include 23 as a source number on this card.</p>			
<p>If do not wish this option, include a blank card for each print interval.</p>			

which the card is read. "Card number" denotes the order within the group (Input Concerning the Differential Equations, General Information for Chain of Radionuclides, etc.) or quantity of cards for each group to be input. "Columns" refers to the actual columns on the card in which the data must be punched. "Remarks, Variables, Etc." gives an explanation as to the variable name, definition, and the form of the data to be punched on the cards. The `FORMAT` descriptor indicates the content and length of the fields on the card. In the `FORMAT` descriptor the letter "A" denotes alphabetic data; "I" refers to integer data (no decimal point) which must always be right-justified in the field; "F" or "E" indicates a real number with or without a signed exponent (decimal point should be punched). The variable names are those used in the `FORTRAN` program. Several examples of input are included in Appendix B.

Output

Output may be grouped in four categories: (1) line printer output on unit 6, (2) line printer output on unit 10, (3) disk output on unit 16, and (4) punched output on unit 7. An explanation of each is given below.

1. Printout on unit 6 consists of the following:
 - a. Summary printout of input - printed in `MAIN`.
 - b. Error messages concerning incorrect input - `MAIN`.
 - c. Coefficient matrix printed by rows - `MATRIX`.
 - d. The nonzero elements of the coefficient matrix and the corresponding row and column index for each nonzero entry - `MATRIX`.

- e. Output for each print interval consisting of possible error messages from DRIVES and value of T and the vector Y printed in YNA.
2. Printout on unit 10 consists of the following:
 - a. Summary of percent deposition - printed in MAIN.
 - b. Listing of the isotope, half life, class, absorption coefficient and branching information for each nuclide - MAIN.
 - c. Retention information for each organ and for each member of the chain - ORGAN.
 - d. For each time period YNA prints the value of T, the cumulated activity ($\mu\text{Ci-days}$), and activity (μCi) entering blood and GIT for each organ and for each member of the chain. OTHER prints the result of the "other tissues" transformation if it is requested.

Printout is divided into two output units so that either unit 6 or unit 10 output may be suppressed. We commonly suppress unit 6 output unless the program abnormally terminates after which unit 6 output should be printed in an additional run for diagnostic purposes.

3. Disk output on unit 16 may be passed to another job step for use in dose calculations as in Eq. (1.1). The unit 16 data set is written unformatted on a 2314 disk pack, with a record format of variable blocked spanned, a logical record length of 150, and a block-size of 1504. The temporary data set is given the name &&UCIDAY and

is created with the disposition of (NEW,PASS). See Appendix C for the summary of Job Control Language (JCL) needed to execute the code. Output on 16 is explained in Table 5.2. In Table 5.2, "Subroutine" refers to the subroutine in which the record is produced. "Record Number" denotes the order in which the records are created. "Remarks, Variables, Etc." gives an explanation of the variable name. "Internal Form of Data" gives the form of the internal data storage. "Alphameric", "integer", and "real" refer to internal computer representation of data.

4. Punched output to unit 7 is performed in routine YNA. For each nuclide in the chain one or more cards are punched which contains the cumulated activity ($\mu\text{Ci-days}$) in the order stated in ISORS (see Table 5.2). These cards are punched with the format (1P8E10.3).

Table 5.2. Form of Data Set Written on Unit 16

Subroutine	Record Number	Remarks, Variables, Etc.	Internal Form of Data
MATRIX	1	NISØ - number of radionuclides in the chain.	integer
		NØRG - number of organs (excluding lung and GIT).	integer
		IØPT - mode of intake. 0 - inhalation 1 - ingestion 2 - injection	integer
If IØPT=1, read record 2.			
MAIN	2	NØRAL - number of oral intakes (for ICRP tables only).	integer
MAIN	3	TAG(ID), ID=1,...,NISØ - atomic symbol and atomic number for each member of the chain.	alphanumeric
MAIN	4	ISØRS(ISØ), ISØ=1,...,total number of organs - indices for organs in the order NP, TB, P, L, S, SI, UL, LL, other organs in their order of input.	integer
For each time period read:			
YNA	1	TIME(ITM) - right end point of the time interval.	real
For each radionuclide read:			
YNA	1	VECTOR(ISØ), ISØ=1,...,total number of organs - cumulated activity given as microcurie-days for each of the organs in the order stated in ISØRS.	real
If IØPT=0 read:			
YNA	2	F(I), I=1,2,3. F(1) - microcuries entering blood from lungs. F(2) - microcuries entering GIT from lungs. F(3) - microcuries entering blood from GIT.	real

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APPENDIX A
PROGRAM LISTING
(Microfiche)

```

C                                     10
C   MAIN DRIVER                                     20
C                                     30
C   IMPLICIT REAL*8 (A-H,O-Z)                                     40
C   COMMON/PPRINT/TPRL (10),DTPR (10),TPRINT,I PRT, NPR, NPRT, NABPRT(10) 50
C   COMMON/SC1275/T, H, HMIN, HMAX, EPS1, NP1, KFLAG1, JSTART, N, NIUV, HHC. 60
C   YRAY(400), ERROR(400), PSAVE(800), YPRINT(400)                                     61
C *** NEED DIMENSION YRAY(NIO), ERROR(NIO), PSAVE(2*NIO) 70
C   COMMON/NSW/NSW                                     80
C   DATA NPRTA/10/                                     90
C   DATA MAXNO/400/                                    100
C *** MUST HAVE NO.LE.MAXNO                                    110
C   DATA NELY/200/                                    120
C   DIMENSION Y(2*00)                                    130
C *** DIMENSION Y(NELY)                                       140
C                                     150
C   NIO=MAXNO                                           160
C ***                                                     170
C ***                                                     180
C *** SET NP AND T1 INTERNALLY                               190
C *** HAVE DEFAULT VALUES FOR N1 AND EPS                   200
C ***                                                     210
C ***                                                     220
C *** DESIRE TO INTEGRATE FROM T1 TO TLAST                 230
C                                     240
C *** NP=21 INDICATES A STIFF SYSTEM.                       250
C   NP=21                                               260
C                                     270
C *** INTEGRATE  $Y'(T) = A * Y(T) + X0$  FROM T=T1 TO T=TLAST WHERE Y(T1)=0. 280
C *** Y IS THE TIME INTEGRAL OF ACTIVITY                   290
C *** A IS A CONSTANT LOWER TRIANGULAR MATRIX AND X0 IS A CONSTANT VECT. 300
C *** A, X0 ARE OBTAINED FROM SUBR. MATRIX BY A CALL IN SUBR.STIFF 310
C   T1=0.00                                             320
C                                     330
C   10 READ(5,1001,END=75) NO,TLAST,N1,EPS              340
C   1001 FORMAT (I5,7E10.2)                             350
C   NSW=1                                                360
C   IF (EPS.EQ.0.00) EPS=1.D-6                          370
C   IF (N1.EQ.0.00) N1=1.D-7                            380
C                                     390
C *** SET INITIAL VALUES TO 0.                            400
C                                     410
C   15 DO 20 I = 1, NO                                    420
C   20 Y(I)=0.00                                         430
C                                     440
C                                     450
C   READ (5,1002) NPRT                                    460
C   1002 FORMAT(I5)                                       470
C   NPREAD=NINO (NPRT, NPRTA)                             480
C   25 READ (5,1003) (TPRL(I),DTPR(I),NABPRT(I),I=1,NREAD) 490
C   1003 FORMAT (2E10.2,I5)                               500
C   PRINT 1004, NPRT, NREAD, (TPRL(I),DTPR(I),NABPRT(I),I=1,NREAD) 510
C   1004 FORMAT( '1NO. PRINT INTERVALS DESIRED = 'I2,3X,'NO. READ = 'I2/ 520
C   '1' POINTS SELECTED IN PRINT INTERVAL = T+Q*DT WHERE Q=0,1,....,N' / 521
C   29X, 'T'18X'DT'11X'N'/(1P19.10,219.10,I6)) 522
C                                     530

```

	IF (NPRD.PQ.NPRT) GO TO 30	540
	PRINT 1005	550
1005	FORMAT(///'O NEED MORE STORAGE SPACE FOR INTERVALS IN MAIN.'//)	560
	GO TO 50	570
	CONTINUE	580
30	THAX=TPRL(NPRT) + NARPT(NPRT)*UTPR(NPRT)	590
		600
C		610
C ***	EXPECT THAX TO BE THE LARGEST T VALUE AT WHICH A 'CALL YOUT' IS	620
C ***	DESTROYED.	630
	IF(DABS(TLAST-THAX).LE.1.0-10*DABS(TLAST)) TLAST=THAX	640
	IF(TLAST.LE.THAX) GO TO 35	650
C		660
	PRINT 1006,TLAST,THAX	670
1006	FORMAT(///'O VALUE OF TLAST HAS BEEN CHANGED FROM'1PE21.13,1X,'TO'	680
	1E21.13//)	681
C		690
	TLAST=THAX	700
35	NT=NPRD-1	710
	T0 = T1	720
	NC=0	730
	IF(NT.LE.1) GO TO 45	740
	DO 40 I=1,NT	750
	IF(TPRL(I)+NARPT(I)*UTPR(I).LT.TPRL(I+1)) GO TO 40	760
	NC=NC+1	770
40	CONTINUE	780
C		790
	IF(NC.GT.0) PRINT 1007,NC,NT	800
1007	FORMAT('1'12,1X'OP THE '12,1X' CONSECUTIVE INTERVAL PAIRS OVERLAP.'	810
	1' PPOG. WILL CALL EXIT. SEE MAIN DRIVER.'/)	811
C		820
45	CONTINUE	830
	IF(T0.LE.TPRL(1)) GO TO 55	840
	PRINT 1008	850
1008	FORMAT('1TPRL(1).LT.T0. CALL EXIT. SEE MAIN DRIVER.')	860
50	CALL EXIT	870
C		880
55	IF(NC.GT.0) GO TO 50	890
	T*PRINT=TPRL(1)	900
	IPRT=1	910
	NPR=0	920
	NCASE=0	930
	NO = N1	940
	WRITE(6,1009) NO, NP, T0, TLAST, NO, EPS	950
1009	FORMAT('0NO='14/' NP='12/' T0='1PE22.15/' TLAST='1E22.15/	960
	1' NO='1E22.15/' EPS='1E10.4/)	961
C		970
C ***	FOR STIPP PROCEDURE NEED 6*NO ELEMENTS IN 'VECTOR' Y. FOR ADAM'S	980
C ***	(NONSTIPP) NEED 13*NO ELEMENTS.	990
C ***	NP=10*NETH+METH, METH=1 FOR ADAM'S, =2 FOR STIPP PROCEDURE.	1000
C		1010
	IF(NP.GT.19) GO TO 60	1020
C		1030
C ***	ADAM'S (NONSTIPP)	1040
C		1050
	IF(13*NO.LE.NELY) GO TO 65	1060


```

      PRINT 1010,NELY                                1070
1010 FORMAT('I NEED 13*NO ELEMENTS IN VECTOR Y IN MAIN. HAVE ONLY NELY='
1I6)                                                1080
      GO TO 50                                        1090
C                                                    1100
60  IF(6*NO.LE.NELY) GO TO 65                        1110
C                                                    1120
C *** STIFF PROCEDURE.                               1130
C                                                    1140
      PRINT 1011,NELY                                1150
1011 FORMAT('I NEED 6*NO ELEMENTS IN VECTOR Y IN MAIN. HAVE ONLY NELY='
1I6)                                                1160
      GO TO 50                                        1170
C                                                    1180
65  IF(NO.LE.NAXNO) GO TO 70                        1190
      PRINT 1012,NO,NAXNO                            1200
1012 FORMAT('NO EXCEEDS MAX.NO/' NO='I6/' NAXNO='I6/' INCREASE DIM. I
IN COMMON SC1235.')                                1210
      GO TO 50                                        1220
C                                                    1230
70  CONTINUE                                         1240
C                                                    1250
C                                                    1260
      CALL DRIVES(NO,TO,TLAST,Y,NO,EPS,NP,KFLAG)     1270
      WRITE (6,1013) KFLAG                            1280
1013 FORMAT('KFLAG = ',I6)                          1290
C                                                    1300
      GO TO 10                                        1310
75  STOP                                             1320
C ----- LAST CARD OF MAIN DRIVER -----          1330
      END                                             1340

```

```

SUBROUTINE DRIVES(NO,TO,TLAST,Y,NO,EPS,NP,KFLAG)
IMPLICIT REAL*8 (A-H,O-Z)
COMMON/SC1235/T,H,HNIN,HMAX,EPS1,NP1,KFLAG1,JSTART,B,NIN,NXO,
1YMAX(400),ERROR(N00),PSAVE(800),YPRINT(400)
C *** NEED DIMENSION YMAX(NX0),ERROR(N00),PSAVE(2*NX0)
COMMON/PRINT/TPR(10),DTPR(10),TPRINT,IPRT,NPR,NPRT,NA NPRT(10)
DIMENSION Y(NO,1)
C*****D257 12
C* DRIVES IS DRIVER SUBROUTINE FOR STIFF. SEE WRITEUP FOR FULL DETAILS.D257 13
C* D257 14
C* THE INPUT PARAMETERS ARE. D257 15
C* NO = THE NUMBER OF DIFFERENTIAL EQUATIONS, INITIALLY. D257 17
C* TO = THE INITIAL VALUE OF T, THE INDEPENDENT VARIABLE. D257 18
C* TLAST = THE FINAL VALUE OF T. D257 19
C* Y = AN NO BY LMAX ARRAY FOR THE COMPUTED SOLUTION, THE SAME D257 20

```



```

IP (NO .LE. 0) GO TO 410                                D257 77
IP ((T0-TLAST)*NO .GE. 0.0D0) GO TO 420                D257 78
N=NO
N = 0                                                    D257 80
H = NO                                                  D257 81
NINV=0
NMIN = DABS(NO)                                         D257 84
NMAX = DABS((T-TLAST) * .1D0)                          D257 85
NMIN = DMIN(NMIN,.1D0*NMAX)                             D257 86
EPS1 = EPS                                              D257 97
NP1 = NP                                                D257 98
JSTART=0
KPLAG = 0                                               D257 90
C *** MAY WISH TO PRINT INITIAL VALUES THRU YOUT.
IP(.EQ.TPRINT) CALL YOUT(NO,TLAST,Y,I)
10 CALL STIPP(Y,NO)
NQ = JSTART                                             D257 93
C
KGO = 1 - KPLAG1                                       D257 94
GO TO ( 20,100,200,300,20,350) , NKO                  D257 95
C KPLAG1 = 0, -1, -2, -3, -5                            D257 96
C
20 CONTINUE                                           D257 97
IP(.GE.TPRINT) CALL YOUT(NO,TLAST,Y,NQ)               D257 98
IP ((T-TLAST)*N .LT. 0.0D0) GO TO 10                  D257 99
C*****D257 127
C* THE PROBLEM IS FINISHED. HERE CALL YOUT AND/OR OTHER ROUTINES D257 128
C* TO OUTPUT DESIRED FINAL RESULTS.                    D257 129
C*****D257 130
30 TO 500                                             D257 131
C
100 WRITE (LOUT,105) T                                  D257 132
105 FORMAT (//30H KPLAG = -1 FROM STIPP AT T = ,D25.16/ D257 133
1 39H ERROR TEST FAILED WITH ABS(N) = NMIN/)          D257 134
110 IP (KPLAG .EQ. 10) GO TO 150                       D257 135
KPLAG = KPLAG + 1                                       D257 136
NMIN = NMIN * .1D0                                       D257 137
N = N * .1D0                                             D257 138
WRITE (LOUT,115) N                                       D257 139
115 FORMAT (24H N HAS BEEN REDUCED TO ,D25.16,        D257 140
1 26H AND STEP WILL BE RETRIED//)                      D257 141
JSTART = -1                                             D257 142
GO TO 10                                                D257 143
C
150 WRITE (LOUT,155)                                    D257 144
155 FORMAT (//40H PROBLEM APPEARS UNSOLVABLE WITH GIVEN INPUT//) D257 145
C*****D257 146
C* NMIN HAS BEEN CUT BY 10 ORDERS OF MAGNITUDE WITH NO SUCCESS. D257 147
C* AT THIS POINT, OUTPUT INFORMATION NEEDED FOR DEBUGGING.   D257 148
C*****D257 149
CALL YOUTD(Y,NQ)                                       D257 150
GO TO 500                                               D257 151
C
200 WRITE (LOUT,205) T,N                                D257 152
205 FORMAT (//30H KPLAG = -2 FROM STIPP AT T = ,D25.16,5H N = ,D25.16/ D257 153
1 52H THE REQUESTED ERROR IS SMALLER THAN CAN BE HANDLED//) D257 154

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C* AT THIS POINT, OUTPUT INFORMATION NEEDED FOR DEBUGGING.
CALL YOUTD(Y,NQ)
GO TO 500
D257 159
D257 160
D257 161
D257 162
C
300 WRITE (LOUT,305) T
D257 163
305 FORMAT(/30H KFLAG = -3 FROM STIFF AT T = ,D25.16/
1 45H CONJECTOR CONVERGENCE COULD NOT BE ACHIEVED/)
D257 164
GO TO 110
D257 165
350 WRITE (LOUT,355) T,N
D257 167
355 FORMAT(/30H KFLAG = -5 FROM STIFF AT T = ,D25.16,
1 5H N = ,D25.16/
2 45H N IS SUCH THAT T>N .EQ. T ON MACHINE. /)
D257 168
D257 169
CALL YOUTD(Y,NQ)
D257 170
GO TO 500
D257 171
D257 172
C
400 WRITE (LOUT,405)
D257 173
405 FORMAT(/20H ILLEGAL INPUT.. EPS .LE. 0.//)
D257 174
KFLAG = -4
D257 175
D257 176
RETURN
D257 177
C
410 WRITE (LOUT,415)
D257 178
415 FORMAT(/25H ILLEGAL INPUT.. N .LE. 0.//)
D257 179
KFLAG = -4
D257 180
D257 181
RETURN
D257 182
C
420 WRITE (LOUT,425)
D257 183
425 FORMAT(/38H ILLEGAL INPUT.. (T0-TLAST)*N0 .GE. 0.//)
D257 184
KFLAG = -4
D257 185
D257 186
RETURN
D257 187
C
500 KFLAG = KFLAG1
D257 188
T0 = T
D257 189
NQ = N
D257 190
RETURN
D257 191
END
D257 192
D257 194

```

```

SUBROUTINE STIFF(Y,NQ)
IMPLICIT REAL*8(A-H,O-Z)
COMMON/NSW/NSW
COMMON/SC1235/T,N,HNIN,HNAX,EPS,NP,KFLAG,JSTART,N,NIN,NIO,
1YMAX(400),RPROR(400),PSAVE(800),YPRINT(400)
C *** NEED DIMENSION YMAX(NX0),ERROR(NIO),PSAVE(2*NX0)
DI MENSION Y(NQ,13),BL(13),PQ(4)
C*****
C* STIFF PERFORMS ONE STEP OF THE INTEGRATION OF AN INITIAL VALUE
C* PROBLEM FOR A SYSTEM OF ORDINARY DIFFERENTIAL EQUATIONS.
C* COMMUNICATION WITH STIFF IS DONE WITH THE FOLLOWING VARIABLES..
D257 206
D257 207
D257 208
D257 209
D257 210

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C* D257 211
 C* Y AN N0 BY LMAX ARRAY CONTAINING THE DEPENDENT VARIABLES D257 212
 C* AND THEIR SCALED DERIVATIVES. LMAX IS CURRENTLY 13 FOR D257 213
 C* THE ADAMS METHODS AND 6 FOR THE GEAR METHODS. LMAX - 1 D257 214
 C* IS MAXDER, THE MAXIMUM ORDER USED. SEE SUBROUTINE TMSOC. D257 215
 C* Y(I,J+1) CONTAINS THE J-TH DERIVATIVE OF Y(I), SCALED BY D257 216
 C* H**J/FACTORIAL(J). ONLY Y(I), 1 .LE. I .LE. N, NEED BE D257 217
 C* SET BY THE CALLING PROGRAM ON THE FIRST ENTRY. D257 218
 C* IF IT IS DESIRED TO INTERPOLATE TO NON-MESH POINTS, D257 219
 C* THE Y ARRAY CAN BE USED. IF THE CURRENT STEP SIZE D257 220
 C* IS H AND THE VALUE AT T + E IS NEEDED, FORM D257 221
 C* S = E/H, AND THEN COMPUTE D257 222
 C* H*
 C* Y(I) (T+E) = SUM Y(I,J+1)*S**J . D257 223
 C* J=0 D257 224
 C* D257 225
 C* THE Y ARRAY SHOULD NOT BE ALTERED BY THE CALLING PROGRAM. D257 226
 C* WHEN REFERENCING Y AS A 2-DIMENSIONAL ARRAY, USE A D257 227
 C* COLUMN LENGTH OF N0, AS THIS IS THE VALUE USED IN STIFF. D257 228
 C* N THE NUMBER OF FIRST ORDER DIFFERENTIAL EQUATIONS. N D257 229
 C* MAY BE DECREASED ON LATER CALLS IF THE NUMBER OF D257 230
 C* ACTIVE EQUATIONS REDUCES, BUT IT MUST NOT BE D257 231
 C* INCREASED WITHOUT CALLING WITH JSTART = 0. D257 232
 C* NO A CONSTANT INTEGER .GE. N, USED FOR DIMENSIONING PURPOSES. D257 233
 C* NO MUST NOT BE CHANGED WITHOUT SETTING JSTART = 0. D257 234
 C* T THE INDEPENDENT VARIABLE. T IS UPDATED ON EACH STEP TAKEN. D257 235
 C* H THE STEP SIZE TO BE ACCEPTED ON THE NEXT STEP. D257 236
 C* H MAY BE ADJUSTED UP OR DOWN BY THE ROUTINE D257 237
 C* IN ORDER TO ACHIEVE AN ECONOMICAL INTEGRATION. D257 238
 C* HOWEVER, IF THE H PROVIDED BY THE USER DOES D257 239
 C* NOT CAUSE A LARGER ERROR THAN REQUESTED, IT D257 240
 C* WILL BE USED. TO SAVE COMPUTER TIME, THE USER IS D257 241
 C* ADVISED TO USE A FAIRLY SMALL STEP FOR THE FIRST D257 242
 C* CALL. IT WILL BE AUTOMATICALLY INCREASED LATER. D257 243
 C* H CAN BE EITHER POSITIVE OR NEGATIVE, BUT ITS SIGN D257 244
 C* MUST REMAIN CONSTANT THROUGHOUT THE PROBLEM. D257 245
 C* HMIN THE MINIMUM ABSOLUTE VALUE OF THE STEP SIZE THAT WILL BE D257 246
 C* USED FOR THE PROBLEM. ON STARTING THIS MUST BE MUCH D257 247
 C* SMALLER THAN THE AVERAGE ABS (H) EXPECTED, SINCE D257 248
 C* A FIRST ORDER METHOD IS USED INITIALLY. D257 249
 C* HMAX THE MAXIMUM ABSOLUTE VALUE OF THE STEP SIZE THAT WILL D257 250
 C* BE USED FOR THE PROBLEM. D257 251
 C* EPS THE RELATIVE ERROR TEST CONSTANT. SINGLE STEP ERROR D257 252
 C* ESTIMATES DIVIDED BY YMAX(I) MUST BE LESS THAN THIS D257 253
 C* IN THE EUCLIDEAN NORM. THE STEP AND/OR ORDER IS D257 254
 C* ADJUSTED TO ACHIEVE THIS. D257 255
 C* MP THE METHOD FLAG. MP IS A POSITIVE INTEGER WITH D257 256
 C* TWO DECIMAL DIGITS--METH AND MITER (MP=10*METH+MITER). D257 257
 C* METH IS THE BASIC METHOD INDICATOR.. D257 258
 C* METH = 1 MEANS THE ADAMS METHODS. D257 259
 C* METH = 2 MEANS THE STIFF SYSTEM METHODS OF GEAR. D257 260
 C* MITER IS THE ITERATION METHOD INDICATOR.. D257 261
 C* MITER = 0 MEANS FUNCTIONAL ITERATION (NO PARTIAL D257 262
 C* DERIVATIVES NEEDED). D257 263
 C* MITER = 1 MEANS CHORD METHOD WITH ANALYTIC JACOBIAN. D257 264
 C* FOR THIS USER SUPPLIES SUBROUTINE D257 265
 C* PEDERV(N,T,Y,PD,N0). PD IS AN N BY N ARRAY, D257 266

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C*          STORED AS AN M0 BY M0 ARRAY. PD(I,J) IS          D257 267
C*          TO BE SET TO THE PARTIAL DERIVATIVE OF          D257 268
C*          YDOT(I) WITH RESPECT TO Y(J).                   D257 269
C*          NITER = 2 MEANS CHORD METHOD WITH JACOBIAN        D257 270
C*          CALCULATED IN STIPP BY FINITE DIFFERENCES.      D257 271
C*          PEDERV IS NOT CALLED AND A JUMPY CAN BE USED.    D257 272
C*          NITER = 3 MEANS CHORD METHOD WITH JACOBIAN REFLACD D257 273
C*          BY A DIAGONAL APPROXIMATION BASED ON A          D257 274
C*          DIRECTIONAL DERIVATIVE.                          D257 275
C*          MAX AN ARRAY OF M LOCATIONS WHICH CONTAINS THE    D257 276
C*          ABSOLUTE VALUE OF EACH Y SEEN SO FAR.            D257 277
C*          ERROR AN ARRAY OF M ELEMENTS PROPORTIONAL TO THE  D257 278
C*          ONE STEP ERROR IN EACH COMPONENT.                D257 279
C*          KFLAG A COMPLETION CODE WITH THE FOLLOWING MEANINGS.. D257 280
C*          0 THE STEP WAS SUCCESSFUL.                       D257 281
C*          -1 THE REQUESTED ERROR COULD NOT BE ACHIEVED     D257 282
C*            WITH ABS(M) = RMIS.                             D257 283
C*          -2 THE REQUESTED ERROR IS SMALLER THAN CAN       D257 284
C*            BE HANDLED FOR THIS PROBLEM.                   D257 285
C*          -3 CORRECTOR CONVERGENCE COULD NOT BE           D257 286
C*            ACHIEVED FOR ABS(M) .GT. RMIS.                 D257 287
C*          -5 M WAS SUCH THAT T+M .EQ. T ON MACHINE.       D257 288
C*          ON A RETURN WITH KFLAG NEGATIVE, THE VALUES OF Y AND
C*          THE Y ARRAY ARE AS OF THE BEGINNING OF THE LAST
C*          STEP, AND M IS THE LAST STEP SIZE ATTEMPTED.    D257 291
C*          JSTART AN INTEGER USED ON INPUT AND OUTPUT.      D257 292
C*          ON INPUT, IT HAS THE FOLLOWING VALUES AND MEANINGS.. D257 293
C*          0 PERFORM THE FIRST STEP. THIS VALUE ENABLES
C*            THE SUBROUTINE TO INITIALIZE ITSELF.           D257 294
C*          .GT.0 TAKE A NEW STEP CONTINUING FROM THE LAST.  D257 296
C*            ASSUMES THE LAST STEP WAS SUCCESSFUL AND
C*            USER HAS NOT CHANGED ANY PARAMETERS.          D257 297
C*          .LT.0 REPEAT THE LAST STEP WITH A NEW VALUE OF  D257 299
C*            M AND/OR EPS AND/OR NP. THIS MAY BE
C*            EITHER IN REDOING A STEP THAT FAILED, OR
C*            IN CONTINUING FROM A SUCCESSFUL STEP.          D257 302
C*          ON EXIT, JSTART IS SET TO M0. THE CURRENT ORDER OF THE
C*          METHOD. THIS IS ALSO THE ORDER OF THE MAXIMUM DERIVATIVE
C*          AVAILABLE IN THE Y ARRAY. AFTER A SUCCESSFUL STEP,
C*          JSTART NEED NOT BE RESET FOR THE NEXT CALL.      D257 306
C*          M A BLOCK OF LOCATIONS USED FOR PARTIAL DERIVATIVES IF
C*          NITER IS NOT 0. IF NITER IS 1 OR 2, ITS LENGTH MUST
C*          BE AT LEAST M0*(M0+1), FOR THE JACOBIAN AND PIVOT
C*          INFORMATION. IF NITER = 3, ITS LENGTH MUST ONLY BE
C*          AT LEAST M.                                       D257 311
C*          PSAVE A BLOCK OF AT LEAST 2*M0 LOCATIONS FOR TEMPORARY STORAGE.
C*          D257 312
C*          THE PARAMETERS WHICH MUST BE INPUT BY THE USER ARE..
C*          M, M0, T, Y, M, RMIS, RMAX, EPS, NP, JSTART.      D257 314
C*          D257 315
C*          D257 316
C*          ADDITIONAL SUBROUTINES REQUIRED ARE..              D257 317
C*          ISOC(METH,M0,EL,T0,MAXDER) SETS METHOD COEFFICIENTS.
C*          IT IS PROVIDED BY THE PACKAGE.                   D257 318
C*          DIPPON(M,T,Y,YDOT) COMPUTES DY/DX GIVEN T AND Y. D257 320
C*          PEDERV(M,T,Y,PD,M0) COMPUTES PARTIAL DERIVATIVES. IT IS USED
C*          ONLY IF NITER = 1. (SEE BP DESCRIPTION.)          D257 321
C*          D257 322

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C* PHOCED(NO,N,A,IPS,S,IER,NIN); AND WLOS(NO,N,A,D,I,IPS) ARE USED D257 323
C* FOR THE SOLUTION OF SYSTEMS OF LINEAR ALGEBRAIC EQUATIONS. D257 324
C* THEY ARE PROVIDED IN THE PACKAGE, AND USED IF NITER = 1 OR 2. D257 325
C* D257 326
C* THE CALLING PROGRAM MUST CONTAIN THE COMMON DECLARATIONS GIVEN D257 327
C* ABOVE, IN WHICH THE LENGTHS ARE AS GIVEN BY THE DESCRIPTIONS D257 328
C* OF THE VARIABLES ABOVE. THE OPTIONAL ROUTINE DRIVES MAY BE D257 329
C* USED FOR THIS PURPOSE. THE CONSTANT ABOISE IN THE DATA STATEMENT D257 330
C* BELOW SHOULD BE SET TO THE NOISE LEVEL OF THE MACHINE. D257 331
C* D257 332
C* THE ORIGINAL VERSION OF THIS PROGRAM WAS WRITTEN AT LLL BY A. C. D257 333
C* WINDMANSR FOR CDC COMPUTERS. THE CDC VERSION WAS MODIFIED FOR D257 334
C* USE ON VAX COMPUTERS AT ARGONNE IN JUNE, 1973. D257 335
C* D257 336
C* LATEST REVISION SEPTEMBER, 1973 D257 337
C* D257 338
C*****D257 339
C DATA ABOISE / 5.D-16/ D257 340
C* D257 341
C*****D257 342
DIMENSION A(12001),IDIAG(400),XO(400),PW(6000)
DIMENSION NBLK(015)
REAL*8 LR(15)
EQUIVALENCE (A(1),IND(1))
DIMENSION IND(1)
DATA NLR/15/
DATA NUDAA/12001/
DATA NUDPW/6000/
C *** NUDPW = NO. WORDS IN VECTOR PW. REQUIRE NUDPW.GE.NBLK
C *** NUDAA = NO. WORDS IN ARRAY A. SHOULD BE ODD
C *** NXO= NO. ELE. IN XO VECTOR = NO. ELE. IN IDIAG VECTOR.
C *** NLR=NO. WORDS IN ARRAY LR.
IF (NLR.EQ.0) GO TO 1
C ***
NLR=1
C ***
NLR=0
NUDAV=NUDAA/2
INDIA=NUDAV+1
C *** WANT A(1) AND A(INDIA) TO BE LOCATED IN DIFFERENT MEMORY BOXES.
C *** ON THE (M 360/91.
IF (15*(NUDAV/16).EQ.NUDA) INDIA=INDIA+1
C *** XO AND IDIAG WILL EACH HAVE 8 ELEMENTS IN THEM ON RETURN FROM
C *** MATRIX
IF (N.LE.NXO) GO TO 3
PRINT *,N,NXO
4 FORNA? ('IFORMAT 8 SUBR.SIIPP'/' N='I4/' NXO=NAIHO='I4//
1' VECTORS WITH DIM (NXO) NEED TO BE INCREASED TO 8 AT LEAST.'//
2' SET NAIHO IN DATA STATEMENT IN 'NAIHO' TO NEW DIM. VALUE OF NXO
3')
10 CALL EXIT
3 CONTINUE
C *** MATRIX RETURNS THE DIAGONAL OF A SPLIT INTO 8 PART IN THE DIAG.
C *** ELEMENTS IN VECTOR A AND INTO THE LR ARRAY THAT EXPANDS TO
C *** DIAG(LR(1)*I(1),...,LR(NDSUBN)*I(NDSUBN)) WHERE I(J) IS A VECTOR
C *** OF 1'S OF DIMENSION NBLK(J).

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C *** THIS SPLIT IS FOR ANOTHER PROGRAM TO CALC. APPROX. EIGENVECTORS.
C *** THE NO. OF NONZERO OFF-DIAGONAL ELEMENTS IN MATRIX A + B
C *** (N= NUMBER OF DIAG. ENTRIES IN MATRIX A) MUST BE .LE. NWDAY WHERE
C *** NWDAY=MAX. NO. WORDS AVAILABLE FOR THE VECTOR A.
C *** EXPECT SUBR. MATRX TO CHECK THAT AVAILABLE STORAGE IS SUFFICIENT
C *** FOR CURR. PROBLEM
      CALL MATRX(A,NWDAY,A(INDKA),IDIF,XO,NIO,N,LR,NBLK,NLR,NDSUBN)
C *** DIMENSION IDIAG(NIO),XC(NIO),LR(NLR),NBLK(NLR)
C *** NDSUBN=NO. OF DIAG. SUBMATRICES IN ARRAY A.
      IF(NDSUBN.LE.NLR) GO TO 5
      PRINT 6,NDSUBN,NLR
6  FORMAT('1SEE SUBR. STIPP'// 'NEED MORE SPACE IN LR,NBLK. DEFINE NLR
1.'// 'NEED' I3, I4' ELEMENTS'// 'HAVE' I3, I4' ELEMENTS')
      GO TO 10
5  CONTINUE
      NLEA=IDIAG(N)
      IF(NLEA.LE.NWDPN) GO TO 8
      PRINT 9,NLEA,NWDPN
9  FORMAT('1SEE SUBR. STIPP'// 'NEED NLEA=' I4, I4' ENTRIES IN VECTOR PN
1.'// 'HAVE NWDPN=' I4, I4' AVAILABLE.')
      GO TO 10
8  CONTINUE
      NB=0
      NB=INCL+INDKA
      DO 11 I=1,N
      NB=MIN0(NB,IND(NB))
11  NB=NB+6
      NB=NB/(-14)
      IF(NB.LE.352) GO TO 13
      PRINT 12,NB
12  FORMAT('1ASSEMBLER LANGUAGE ROUTINES YPAY AND AYED CAN HANDLE AT M
10ST 352 NONZERO ENTRIES IN A ROW OF THE MATRIX A.'//
2.' MAX. NO. NONZERO ENTRIES IN CURRENT A = NB =' I3)
      GO TO 10
13  CONTINUE
      ICUT=0
      INBLK=0
      DO 20 I=1,N
      IF(I.LE.ICUT) GO TO 21
      INBLK=INBLK+1
      ICUT=ICUT+INBLK(*INBLK)
      RLR=LR(INBLK)
21  CONTINUE
      ID=IDIAG(I)
20  A(ID)=A(ID)+RLR
C *** BASIC DIFFERENTIAL EQ. IS Y'=-A*Y+X0, Y(0)=0
C *** INITIAL CONDITIONS SET IN MAIN PROGRAM.
      NJAC=0.
1  CONTINUE
C*
      KFLAG = 0
      TOLD = T
C* TEST FOR SMALL N ADDED AT ARGONNE IN AUGUST, 1973.
      IF(T+N .EQ. 1) GO TO 685
      IF (JSTART .GT. 0) GO TO 200
      IF (JSTART .NE. 0) GO TO 120

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D257 343
D257 344
D257 345
D257 346
D257 347
D257 348
D257 349

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*****D257 350
C* ON THE FIRST CALL, THE ORDER IS SET TO 1 AND THE INITIAL D257 351
C* DERIVATIVES ARE CALCULATED. YHAR IS INITIALIZED USING THE INITIAL D257 352
C* Y AND YDOT. IF BOTH ARE INITIALLY ZERO IN ANY COMPONENT, THE DEFAULT D257 353
C* VALUE IS 1. RMAX IS THE MAXIMUM RATIO BY WHICH H CAN BE INCREASED D257 354
C* IN A SINGLE STEP. IT IS INITIALLY 1.E4 TO COMPENSATE FOR THE SMALL D257 355
C* INITIAL H, BUT THEN IS NORMALLY EQUAL TO 10. IF A FAILURE D257 356
C* OCCURS (IN CORRECTOR CONVERGENCE OR ERROR TEST), RMAX IS SET AT 2 D257 357
C* FOR THE NEXT INCREASE. EPSJ IS USED AS THE RELATIVE INCREMENT D257 358
C* TO Y WHEN GETTING PARTIALS BY FINITE DIFFERENCING. D257 359
*****D257 360
      NSQ = N0 * N0 D257 361
      NSQ1 = NSQ + 1 D257 362
      N1 = N0 + 1 D257 363
      CALL YFAT(PSAVE,A,Y,X0,A(INDEX),N)
      DO 110 I=1,N
      Y(I,2) = PSAVE(I) * N
      AYI = DABS(Y(I,1))
      IF (AYI.EQ.0.D0) AYI = DABS(Y(I,2))
      IF (AYI.EQ.0.D0) AYI = 1. D0
      YHAR(I) = AYI
110 CONTINUE
      NQ = 1 D257 374
      L = 2 D257 375
      RMAX = 1.0D4 D257 376
      EPSJ = DSQRT(ANOISE) D257 377
      CRATE = 1.0D0 D257 378
      OLDLO = 1.0D0 D257 379
      FC = 0.0D0 D257 380
      YFOLD = 0 D257 381
      NETH = 0 D257 382
      NOLD = N D257 383
C* THIS INITIALIZES NITER SO STIFF WILL FUNCTION CORRECTLY WHEN D257 384
C* GIVEN THE SAME NITER AS WAS GIVEN IN A PREVIOUS PROBLEM. D257 385
      NITER = 0 D257 386
C* ADDED AT ARGONNE IN JULY, 1973. D257 387
*****D257 388
C* IF THE CALLER HAS CHANGED NETH, OR IF JSTART = 0, TESOC IS CALLED D257 389
C* TO SET THE COEFFICIENTS OF THE METHOD. IF THE CALLER HAS CHANGED D257 390
C* EPS OR NETH, THE CONSTANTS E, EDN, EDP, AND BND MUST BE RESET. D257 391
C* E IS A COMPARISON FOR ERRORS OF THE CURRENT ORDER NQ. EUP IS D257 392
C* TO TEST FOR INCREASING THE ORDER, EDN FOR DECREASING THE ORDER. D257 393
C* BND IS USED TO TEST FOR CONVERGENCE OF THE CORRECTOR ITERATES. D257 394
C* IF THE CALLER HAS CHANGED N, Y MUST BE RESCALED. D257 395
C* IF N OR NETH HAS BEEN CHANGED, IDOUB IS RESET TO L + 1 TO PREVENT D257 396
C* FURTHER CHANGES IN N FOR THAT MANY STEPS. ALSO, RC IS RESET. D257 397
C* RC IS THE RATIO OF NEW TO OLD VALUES OF THE COEFFICIENT L(0) * N. D257 398
C* WHEN RC DIFFERS FROM 1 BY MORE THAN 30 PERCENT, OR THE CALLER HAS D257 399
C* CHANGED NITER, INEVAL IS SET TO NITER TO FORCE THE PARTIALS TO BE D257 400
C* UPDATED, IF PARTIALS ARE USED. D257 401
*****D257 402
120 IF (NF.EQ.0FOLD) GO TO 150 D257 403
      NBO = NETH D257 404
      NIO = NITER D257 405
      NETH = NF / 10 D257 406
      NITER = NF - 10 * NETH D257 407

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NPOLD = NP
IF (NITER .NE. NIO) IUEVAL = NITER
IF (NETH .EQ. NEO) GO TO 150
IDOMB = L + 1
IPET = 1
130 CALL PESOC(METH,NQ,EL,TQ,MAXDER)
RC = RC * EL(1) / OLDLO
OLDLO = EL(1)
140 EDN = (TQ(1) * EPS) ** 2
E = (TQ(2) * EPS) ** 2
WEP = (TQ(3) * EPS) ** 2
BND = (TQ(4) * EPS) ** 2
GO TO (160, 170, 200), IPET
150 IF (EPS .EQ. EPSOLD) GO TO 160
IPET = 1
GO TO 140
160 LMAX = MAXDER + 1
EPSOLD = EPS
IF (N .EQ. NOLD) GO TO 200
RN = N / NOLD
N = NOLD
170 RN = DNAXI(RN,HNIN/DABS(N))
RN=DMIN1(RN, RNAXI/DABS(N),RNAXI)
R1 = 1.000
C
DO 190 J = 2, L
R1 = R1 * RN
C
DO 180 I = 3L, N
Y(I,J) = Y(I,J) * R1
180 CONTINUE
C
N = N * RN
RC = RC * RN
IDOMB = L + 1
IF (.NE. TOLD) GO TO 690
C*****
C* THIS SECTION COMPUTES THE PREDICTED VALUES BY EFFECTIVELY
C* MULTIPLYING THE Y ARRAY BY THE PASCAL TRIANGLE MATRIX.
C*****
200 IF (DABS(RC-1.000) .GT. 0.300) IUEVAL = NITER
T = T + N
C
DO 210 J1 = 1, NQ
C
DO 210 J2 = J1, NQ
J = NQ - J2 + J1
C
DO 210 I=NL, N
Y(I,J) = Y(I,J) + Y(I,J+1)
210 CONTINUE
C*****
C* UP TO 3 CORRECTOR ITERATIONS ARE TAKEN. CONVERGENCE IS TESTED
C* BY REQUIRING CHANGES TO BE LESS THAN BND, WHICH IS DEPENDENT ON
C* EPS, IN EUCLIDEAN NORM. THE SUM OF THE CORRECTIONS IS ACCUMULATED
C* IN THE VECTOR ERROR(I). IT IS APPROXIMATELY EQUAL TO THE L-TH

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420 IF (INEVAL .EQ. -1) GO TO 440                                D257 604
    T = TCID                                                    D257 605
    YMAX = 2.000                                                D257 606
C                                                                D257 607
    DO 430 J1 = 1, NQ                                           D257 608
C                                                                D257 609
        DO 430 J2 = J1, NQ                                       D257 610
            Y = NQ - J2 * J1                                       D257 611
C                                                                D257 612
            DO 430 I =NL,N                                         D257 613
                Y(I,J) = Y(I,J) - Y(I,J+1)                       D257 614
430 CONTINUE                                                    D257 615
C                                                                D257 616
    IF (DABS(N) .LE. (NMIN*1.0000100)) GO TO 680                D257 617
    RN=.75DC
    TO TO 170
440 INEVAL = NITER
    TO TO 220
C*****D257 622
C* THE CORRECTOR HAS CONVERGED. INEVAL IS SET TO -1 IF PARTIAL
C* DERIVATIVES WERE USED, TO SIGNAL THAT THEY MAY NEED UPDATING ON
C* SUBSEQUENT STEPS. THE ERROR TEST IS MADE AND CONTROL PASSES TO
C* STATEMENT 500 IF IT FAILS.
C*****D257 623
450 D = 0.001                                                  D257 624
C                                                                D257 625
    DO 460 I =NL,N                                             D257 626
        D = D + (ERROR(I) / YMAX(I)) ** 2                       D257 627
C                                                                D257 628
    IF (NITER .NE. 0) INEVAL = -1                               D257 629
    IF (D .GT. D) GO TO 500                                     D257 630
C                                                                D257 631
C*****D257 632
C* AFTER A SUCCESSFUL STEP, UPDATE THE Y ARRAY AND YMAX.
C* CONSIDER CHANGING N IF IDOUB = 1. OTHERWISE DECREASE IDOUB BY 1.
C* IF IDOUB IS THEN 1 AND NQ .LT. MAXDER, THEN ERROR IS SAVED FOR
C* USE IN A POSSIBLE ORDER INCREASE ON THE NEXT STEP.
C* IF A CHANGE IN N IS CONSIDERED, AN INCREASE OR DECREASE IN ORDER
C* BY ONE IS CONSIDERED ALSO. A CHANGE IN N IS MADE ONLY IF IT IS BY A
C* FACTOR OF AT LEAST 1.1. IF NOT, IDOUB IS SET TO 10 TO PREVENT
C* TESTING FOR THAT MANY STEPS.
C*****D257 633
    RPLAG = 0                                                  D257 634
C                                                                D257 635
    DO 470 J = 1, L                                             D257 636
C                                                                D257 637
        DO 470 I =NL,N                                           D257 638
            Y(I,J) = Y(I,J) + EL(J) * ERROR(I)                 D257 639
470 CONTINUE                                                    D257 640
C                                                                D257 641
    DO 490 I =NL,N                                             D257 642
        YMAX(I) = DMAX1(YMAX(I), DABS(Y(I,T)))                 D257 643
C                                                                D257 644
    IF (IDOUB .EQ. 1) GO TO 520                                  D257 645
    IDOUB = IDOUB - 1                                           D257 646
    IF (IDOUB .GT. 1) GO TO 700                                  D257 647
    IF (NQ .EQ. MAXDER) GO TO 700                               D257 648

```

```

C
DO 490 I = NL, N
490 Y(I, LMAX) = ERROR(I)
C
GO TO 700
C*****
C* THE ERROR TEST FAILED. KFLAG KEEPS TRACK OF MULTIPLE FAILURES.
C* RESTORE I AND THE Y ARRAY TO THEIR PREVIOUS VALUES, AND PREPARE
C* TO TRY THE STEP AGAIN. COMPUTE THE OPTIMUM STEP SIZE FOR THIS OR
C* ONE LOWER ORDER.
C*****
500 KFLAG = KFLAG - 1
I = "CID
C
DO 510 J1 = 1, NQ
C
DO 510 J2 = J1, NQ
J = NQ - J2 + J1
C
DO 510 I = NL, N
Y(I, J) = Y(I, J) - Y(I, J+1)
510 CONTINUE
C
RMAX = 2.000
IF (DABS(R) .LE. (RMIN*1.00001D0)) GO TO 660
IF (KFLAG .LE. -3) GO TO 640
PR3 = 1.0D20
GO TO 540
C*****
C* REGARDLESS OF THE SUCCESS OR FAILURE OF THE STEP, FACTORS
C* PR1, PR2, AND PR3 ARE COMPUTED, BY WHICH H COULD BE DIVIDED
C* AT ORDER NQ - 1, ORDER NQ, OR ORDER NQ + 1, RESPECTIVELY.
C* IN THE CASE OF FAILURE, PR3 = 1.0D20 TO AVOID AN ORDER INCREASE.
C* THE SMALLEST OF THESE IS DETERMINED AND THE NEW ORDER CHOSEN
C* ACCORDINGLY. IF THE ORDER IS TO BE INCREASED, WE COMPUTE ONE
C* ADDITIONAL SCALED DERIVATIVE.
C*****
520 PR3 = 1.0D20
IF (NQ .EQ. NAXDEM) GO TO 540
D1 = 0.0D0
C
DO 530 I = NL, N
530 D1 = D1 + ((ERROR(I) - Y(I, LMAX)) / YMAX(I)) ** 2
C
ENQ3 = 0.5D0 / DFLOAT(L+1)
PR3 = ((D1 / ENQ3) ** ENQ3) * 1.4D0 + 1.4D-06
540 ENQ2 = 0.5D0 / DFLOAT(L)
PR2 = ((D / ENQ2) ** ENQ2) * 1.2D0 + 1.2D-06
PR1 = 1.0D20
IF (NQ .EQ. 1) GO TO 560
D = 0.0D0
C
DO 550 I = NL, N
550 D = D + (Y(I, L) / YMAX(I)) ** 2
C
ENQ1 = 0.5D0 / DFLOAT(NQ)

```

```

PR1 = ((D / 2DN) ** ENQ1) * 1.300 * 1.3D-06
560 IF (PR2 .LT. PF3) GO TO 570
    IF (PR3 .LT. PF1) GO TO 590
570 IF (PR2 .GT. PF1) GO TO 580
    NEWQ = NQ
    RH = 1.000 / PR2
    GO TO 620
590 NEWQ = NQ - 1
    C   IUEVAL=4ITER
    RH = 1.000 / PR1
    GO TO 620
590 NEWQ = L
    RH = 1.000 / PR3
    IF (RH .LT. 1.100) GO TO 610
C
    DO 630 I =NL, N
600   Y(I, NEWQ+1) = ERROR(I) * E1(L) / DFLOAT(L)
C
    TO TO 630
610 IDOOR = 10
    GO TO 700
620 IF ((KFLAG .EQ. 0). AND. (RH .LT. 1.100)) GO TO 610
C*****
C* IF THERE IS A CHANGE OF ORDER, RESET NQ, L, AND THE COEFFICIENTS.
C* IN ANY CASE H IS RESET ACCORDING TO RH AND THE Y ARRAY IS RESCALED.
C* THEN EXIT FROM 690 IF THE STEP WAS OK, OR RTDC THE STEP OTHERWISE.
C*****
    IF (NEWQ .EQ. NQ) GO TO 170
630 NQ = NEWQ
    L = NQ + 1
    IRET = 2
    GO TO 130
C*****
C* CONTROL REACHES THIS SECTION IF 3 OR MORE FAILURES HAVE OCCURED.
C* IT IS ASSUMED THAT THE DERIVATIVES THAT HAVE ACCUMULATED IN THE
C* Y ARRAY HAVE ERRORS OF THE WRONG ORDER. HENCE THE FIRST
C* DERIVATIVE IS RECOMPUTED, AND THE ORDER IS SET TO 1. THEN
C* H IS REDUCED BY A FACTOR OF 10, AND THE STEP IS RETRIED.
C* AFTER A TOTAL OF 7 FAILURES, AN EXIT IS TAKEN WITH KFLAG = -2.
C*****
640 IF (KFLAG .EQ. -7) GO TO 670
    RH = .100
    RH = DMAX1(RHMIN/DABS(N), RH)
    H = H * RH
    CALL YPAY(PSAVE, A, Y, X0, A(INDXA), N)
C
    DO 650 I =NL, N
650   Y(I, 2) = H * PSAVE(I)
C
    IUEVAL  NITER
    IDOOR = 10
    IF (NQ .EQ. 1) GO TO 200
    NQ = 1
    L = 2
    IRET = 1
    GO TO 130

```

D257 716
D257 717
D257 718
D257 719
D257 720
D257 721
D257 722
D257 723
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D257 727
D257 728
D257 729
D257 730
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D257 757
D257 758
D257 759
D257 761
D257 762
D257 763
D257 764
D257 765
D257 766
D257 767
D257 768
D257 769
D257 770
D257 771

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C*****D257 772
C* ALL RETURNS ARE MADE THROUGH THIS SECTION. N IS SAVED IN NOLD D257 773
C* TO ALLOW THE CALLER TO CHANGE N ON THE NEXT STEP. D257 774
C*****D257 775
660 KPLAG = -1 D257 776
    GO TO 700 D257 777
670 KFLAG = -2 D257 778
    GO TO 700 D257 779
680 KFLAG = -3 D257 780
    GO TO 700 D257 781
685 KFLAG = -5 D257 782
    GO TO 700 D257 783
690 NNEW = 10.000 D257 784
700 NOLD = N D257 785
    JSTART = N0 D257 786
    RETURN D257 787
C***** LAST CARD OF STIFF *****D257 788
END D257 789

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SUBROUTINE TESOC(NETH, NQ, EL, TQ, NAXDER) D257 793
C* D257 794
C
C 9/ 5/7400
C
    IMPLICIT REAL*8(A-H,O-Z)
    DIMENSION PERTST(12,2,3), EL(13), TQ(3) D257 796
C*****D257 797
C* TESOC IS CALLED BY STIFF AND SETS COEFFICIENTS FOR USE THERE. D257 798
C* THE VECTOR EL, OF LENGTH NQ + 1, DETERMINES THE BASIC METHOD. D257 799
C* THE VECTOR TQ, OF LENGTH 3, IS INVOLVED IN ADJUSTING THE STEP SIZE D257 800
C* IN RELATION TO TRUNCATION ERROR. ITS VALUES ARE GIVEN BY THE D257 801
C* PERTST ARRAY. D257 802
C* THE VECTORS EL AND TQ DEPEND ON NETH AND NQ. D257 803
C* TESOC ALSO SETS NAXDER, THE MAXIMUM ORDER OF THE METHOD AVAILABLE. D257 804
C* CURRENTLY IT IS 12 FOR THE ADAMS METHODS AND 5 FOR THE GEAR METHODS. D257 805
C* LNDI = NAXDER + 1 IS THE NUMBER OF COLUMNS IN THE Y ARRAY. D257 806
C* THE MAXIMUM ORDER USED MAY BE REDUCED SIMPLY BY CHANGING THE D257 807
C* THE NUMBERS IN STATEMENTS 1 AND 2 BELOW. D257 808
C* D257 809
C* THE COEFFICIENTS IN PERTST NEED BE GIVEN TO ONLY ABOUT D257 810
C* ONE PERCENT ACCURACY. THE ORDER IN WHICH THE GROUPS APPEAR BELOW D257 811
C* IS.. COEFFICIENTS FOR ORDER NQ - 1, COEFFICIENTS FOR ORDER NQ, D257 812
C* COEFFICIENTS FOR ORDER NQ + 1. WITHIN EACH GROUP ARE THE D257 813
C* COEFFICIENTS FOR THE ADAMS METHODS, FOLLOWED BY THOSE FOR THE D257 814
C* GEAR METHODS. D257 815
C* D257 816
C* THE ORIGINAL VERSION OF THIS PROGRAM WAS WRITTEN AT LLL BY A. C. D257 817

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C* BINDER FOR CDC COMPUTERS. THE CDC VERSION WAS MODIFIED FOR      D257 818
C* USE ON IBM COMPUTERS AT ARGONNE IN JUNE, 1973.                  D257 819
C*                                                                     D257 820
C* LATEST REVISION SEPTEMBER, 1973                                D257 821
C*                                                                     D257 822
C*****D257 823
DATA BEETS / 1.000,1.000,2.000,1.000,,.315800,.0740700,.0139100,
1 .00218200,.000294500,.0000349200,.00000369200,.000000352400, D257 824
1 1.00,1.00,.500,.166700,.0816700,1.00,1.00,1.00,1.00,1.00,1.00, D257 825
1 1.00,2.00,12.00,24.00,37.8900,53.3300,70.0800,87.9700,106.900, D257 827
1 126.700,147.400,168.800,191.000, D257 828
1 2.000,4.500,7.33300,10.4200,13.700,1.00,1.00,1.00,1.00,1.00,1.00, D257 829
1 1.000,12.000,24.000,37.8900,53.3300,70.0800,87.9700,106.900, D257 830
1 126.700,147.400,168.800,191.000,1.00,3.000, D257 831
1 6.000,3.16700,12.500,1.00,1.00,1.00,1.00,1.00,1.00,1.00,1.000/ D257 832
C*                                                                     D257 833
C*****D257 834
C*                                                                     D257 835
      PL(2) = 1.000 D257 836
      GO TO (1,2), NETA D257 837
1     NAXDER = 12 D257 838
      GO TO (101,102,103,104,105,106,107,108,109,110,111,112), NQ D257 839
2     NAXDER = 5 D257 840
      GO TO (201,202,203,204,205), NQ D257 841
C*****D257 842
C* THE FOLLOWING COEFFICIENTS SHOULD BE DEFINED TO D257 843
C* MACHINE ACCURACY. FOR EACH ORDER NQ, THEY CAN BE CALCULATED D257 844
C* FROM THE GENERATING POLYNOMIAL, D257 845
C*  $L(T) = PL(1) + PL(2)T + \dots + PL(NQ+1)T^{NQ}$ . D257 846
C* FOR THE IMPLICIT ADAMS METHODS, L(T) IS GIVEN BY D257 847
C*  $DL/DT = (T+1)(T+2) \dots (T+NQ-1)/K$ ,  $L(-1) = 0$ , D257 848
C* WHERE  $K = \text{FACTORIAL}(NQ-1)$ . D257 849
C* FOR THE GEAR METHODS, D257 850
C*  $L(T) = (T+1)(T+2) \dots (T+NQ)/K$ , D257 851
C* WHERE  $K = \text{FACTORIAL}(NQ) * (1 + 1/2 + \dots + 1/NQ)$ . D257 852
C* D257 853
C* THE ORDER IN WHICH THE GROUPS APPEAR BELOW IS.. D257 854
C* IMPLICIT ADAMS METHODS OF ORDERS 1 TO 12, D257 855
C* STIFFLY STABLE GEAR METHODS OF ORDERS 1 TO 5. D257 856
C*****D257 857
101  PL(1) = 1.000 D257 858
      GO TO 900 D257 859
102  PL(1) = 0.500 D257 860
      PL(3) = 0.500 D257 861
      GO TO 900 D257 862
103  PL(1) = 4.1666666666666667D-01 D257 863
      PL(3) = 0.7500 D257 864
      PL(4) = 1.6666666666666667D-01 D257 865
      GO TO 900 D257 866
104  PL(1) = 0.17500 D257 867
      PL(2) = 9.1666666666666667D-01 D257 868
      PL(4) = 3.3333333333333333D-01 D257 869
      PL(5) = 4.1666666666666667D-02 D257 870
      GO TO 900 D257 871
105  PL(1) = 3.4861111111111111D-01 D257 872
      PL(3) = 1.0416666666666667D0 D257 873

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	EL (4) = 4.861111111111111D-01	D257 874
	EL (5) = 1.0416666666666667D-01	D257 875
	EL (6) = 9.333333333333333D-03	D257 876
	GO TO 900	D257 877
105	EL (1) = 3.298611111111111D-01	D257 878
	EL (2) = 1.1416666666666667D+00	D257 879
	EL (3) = 0.425D+00	D257 880
	EL (4) = 1.773333333333333D-01	D257 881
	EL (5) = 0.025D+00	D257 882
	EL (6) = 1.388888888888889D-03	D257 883
	GO TO 900	D257 884
107	EL (1) = 3.1559193121693122D-01	D257 885
	EL (3) = 1.225D+00	D257 886
	EL (4) = 7.518518518518518D-01	D257 887
	EL (5) = 2.552093333333333D-01	D257 888
	EL (6) = 4.861111111111111D-02	D257 889
	EL (7) = 4.851111111111111D-03	D257 890
	EL (8) = 1.5341269941269941D-04	D257 891
	GO TO 900	D257 892
108	EL (1) = 3.0422453703703704D-01	D257 893
	EL (2) = 1.2164285714285714D+00	D257 894
	EL (3) = 9.665185185185185D-01	D257 895
	EL (4) = 2.357638888888889D-01	D257 896
	EL (5) = 7.777777777777777D-02	D257 897
	EL (6) = 1.0549148148148148D-02	D257 898
	EL (7) = 7.9355079365079365D-04	D257 899
	EL (8) = 2.4301587301587302D-05	D257 900
	GO TO 900	D257 901
109	EL (1) = 2.9486800044091711D-01	D257 902
	EL (3) = 1.3589285714285714D+00	D257 903
	EL (4) = 9.7655423280423280D-01	D257 904
	EL (5) = 0.4171875D+00	D257 905
	EL (6) = 1.1135416666666667D-01	D257 906
	EL (7) = 0.01875D+00	D257 907
	EL (8) = 1.9345238095238095D-03	D257 908
	EL (9) = 1.1160714285714286D-04	D257 909
	EL (10) = 2.7557319223985891D-06	D257 910
	GO TO 900	D257 911
110	EL (1) = 2.8697544642857143D-01	D257 912
	EL (3) = 1.4144841269841270D+00	D257 913
	EL (4) = 1.0772156084656085D+00	D257 914
	EL (5) = 4.9856701940035273D-01	D257 915
	EL (6) = 0.1484375D+00	D257 916
	EL (7) = 2.9060570987654321D-02	D257 917
	EL (8) = 3.7202380952380952D-03	D257 918
	EL (9) = 2.9964584656084655D-04	D257 919
	EL (10) = 1.3778659611992945D-05	D257 920
	EL (11) = 2.7557319223985891D-07	D257 921
	GO TO 900	D257 922
111	EL (1) = 2.8018959644393672D-01	D257 923
	EL (3) = 1.4644841269841270D+00	D257 924
	EL (4) = 1.1715145502645503D+00	D257 925
	EL (5) = 5.7935819003527337D-01	D257 926
	EL (6) = 1.8832286155202822D-01	D257 927
	EL (7) = 4.1430362654320988D-02	D257 928
	EL (8) = 6.2111441798941799D-03	D257 929

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PL (9) = 6.25206679894179890-04
PL (10) = 4.04174015235126400-05
PL (11) = 1.51565255731922400-06
PL (12) = 2.50521093854417190-08
GO TO 900
112 PL (1) = 2.74265540031599060-01
    PL (3) = 1.50393867243867240+00
    PL (4) = 1.26027116402116400+00
    PL (5) = 6.59234192098765430-01
    PL (6) = 2.30458002645502650-01
    PL (7) = 5.56972461052322160-02
    PL (8) = 9.43948412599412700-03
    PL (9) = 1.11927496693121690-03
    PL (10) = 9.09391534391534390-05
    PL (11) = 4.82253096419753090-06
    PL (12) = 1.50312650312450310-07
    PL (13) = 2.04767569878680990-09
GO TO 900
201 PL (1) = 1.00+00
GO TO 900
202 PL (1) = 6.66666666666666670-01
    PL (3) = 2.33333333333333330-01
GO TO 900
203 PL (1) = 5.45454545454545450-01
    PL (3) = PL (1)
    PL (4) = 9.09090909090909090-02
GO TO 900
204 PL (1) = 0.490+00
    PL (3) = 0.70+00
    PL (4) = 0.20+00
    PL (5) = 0.020+00
GO TO 900
205 PL (1) = 4.37956204379562040-01
    PL (3) = 8.21167893211678830-01
    PL (4) = 3.10218978102189780-01
    PL (5) = 5.47445255474452550-02
    PL (6) = 3.64963503649635040-03
C*
900 DO 910 K = 1, 3
910   TQ(K) = PERTST(NQ, NBTN, K)
C*
    TQ(4) = 0.500 * TQ(2) / DPLOAT(NQ + 2)
    RETURN
C*****LAST CARD OF TESSC*****
END

```

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0257 930
0257 931
0257 932
0257 933
0257 934
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0257 969
0257 970

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C		RATR 50
C	CINDEX(1510)	RATR 70
C	CINDEX(I) = COLUMN INDEX OF A(I)	RATR 80
C		RATR 90
C	ICLASS(10)	RATR 100
C	ICLASS(I) = CLASS OF THE ITH MEMBER OF CHAIN	RATR 110
C		RATR 120
C	ICON(10,10)	RATR 130
C	ICON(I,J) = NO. OF COMPARTMENT FOR DAUGHTER I, ORGAN J	RATR 140
C		RATR 150
C	INDXA()	RATR 160
C	INDXA(2,2*I-1) = -10*(NO. OF NON-ZERO ELEMENTS IN	RATR 170
C	ROW I OF A	RATR 180
C	INDXA(2,2*I) = 10*(NO. OF NON-ZERO ELEMENTS IN	RATR 190
C	ROW I OF A	RATR 200
C	INDXA(1,J) = 10*(J-1) IFF A(I) BELONGS IN COLUMN	RATR 210
C	J OF THE MATRIX A	RATR 220
C		RATR 230
C	ISORS(20)	RATR 240
C	ISORS(I) = INDEX OF THE ITH SOURCE ORGAN (INDICES FROM SEE CODE)	RATR 250
C		RATR 260
C	LAMB(10,4)	RATR 270
C	LAMB(I,1) = LAMBDA OF ABSORPTION IN S FOR DAUGHTER I	RATR 280
C	LAMB(I,2) = LAMBDA OF ABSORPTION IN SI FOR DAUGHTER I	RATR 290
C	LAMB(I,3) = LAMBDA OF ABSORPTION IN OLI FOR DAUGHTER I	RATR 300
C	LAMB(I,4) = LAMBDA OF ABSORPTION IN LII FOR DAUGHTER I	RATR 310
C		RATR 320
C	LAMR(10)	RATR 330
C	LAMR(I) = LN 2/(HALPLIFE OF DAUGHTER I)	RATR 340
C		RATR 350
C	LAMX(10,10)	RATR 360
C	LAMX(I,J) = LN 2/(HALPLIFE OF DAUGHTER I IN COMPARTMENT J)	RATR 370
C		RATR 380
C	LP(K) = LAMR(K)	RATR 390
C		RATR 400
C	NBLK(MDSUBM)	RATR 410
C	NBLK(I) = ORDER OF SQUARE DIAGONAL SUBMATRIX I OF A	RATR 420
C	NBLK(I) = NCON(I)	RATR 430
C		RATR 440
C	NCON(10)	RATR 450
C	NCON(I) = NO. OF COMPARTMENTS FOR DAUGHTER I	RATR 460
C		RATR 470
C	ORGN(10)	RATR 480
C	ORGN(I) = NAME OF THE ITH ORGAN	RATR 490
C		RATR 500
C	PRNT(1000)	RATR 510
C	PRNT(I) = ITH ENTRY OF A GIVEN ROW OF THE COEFFICIENT MATRIX	RATR 520
C		RATR 530
C	RINDEX(1510)	RATR 540
C	RINDEX(I) = ROW INDEX OF A(I)	RATR 550
C		RATR 560
C	TAG(10)	RATR 570
C	TAG(I) = ATOMIC SYMBOL AND ATOMIC NUMBER FOR DAUGHTER I	RATR 580
C		RATR 590
C	TR(10)	RATR 600
C	TR(I) = HALPLIFE OF DAUGHTER I	RATR 610


```

C
C
C      IMPLICIT REAL*8 (A-H,O-Z)
C      REAL*8 LR(1),LANX(10,10),LANR(10),LANAB(10,4)
C      INTEGER*2 RINDEX(12001),CINDEX(12001),SUN1
C      DIMENSION A(1),XO(1),INDXA(2,1),IDTAG(1),ICLASS(10),TR(10),
C      1 PRNT(400),NBLK(1),ISORS(20),P(10),NBRNCH(10),IBRNCH(10)
C
C      COMMON/TABLES/ LANAB(10,4),LANX(10,10),ORGNR(10),TAG(10),
C      1 ICON(10,10),IOPT,NCON(10),NISO,NORG
C      COMMON/SWITCH/ ISWTR
C
C      DATA PLN2/.6931471905599450/
C      DATA ISORS/9,9,10,11,2,3,4,5/
C
C
C      READ NISO = NO. OF RADIONUCLIDES IN CHAIN
C      D3, D4, D5 = REGIONAL DEPOSITION VALUES
C      NORG = NO. OF ORGANS (EXCLUDING LUNG AND GI
C      IOPT = 0 INTAKE BY INHALATION
C      1 INTAKE BY INGESTION
C      2 INTAKE BY INJECTION
C
C      READ(50,1001) NISO, D3, D4, D5, NORG, IOPT, NORAL
C      1001 FORMAT(I10,3F10.3,3I10)
C      WRITE(10,1002)
C      1002 FORMAT(' ')
C      IF(IOPT.EQ.0) WRITE(10,1003) D3, D4, D5
C      1003 FORMAT(' PERCENT DEPOSITION'/' ',4X,'N-P',12X,'T-B',
C      1 12X,'P'/' ',P7.2,4(8X,P7.2),/)
C      WRITE(16) NISO,NORG,IOPT
C      IF(IOPT.EQ.1) WRITE(16) NORAL
C
C      ASSIGN REGIONAL DEPOSITION FOR EACH PATHWAY
C
C      IF(IOPT.EQ.0) CALL LUNG(D3,D4,D5)
C
C      FOR EACH RADIONUCLIDE READ
C      TAG = ATOMIC NO. AND SYMBOL
C      TR = HALFLIFE
C      ICLASS = CLASS
C      NIAB = 0 ABSORPTION IN SI ONLY
C      1 ABSORPTION IN ALL REGIONS OF GI TRACT
C
C      WRITE(10,1004)
C      1004 FORMAT(' ISOTOPE',4X,'HALFLIFE',4X,'CLASS',3X,'LANAB',
C      1 3X,'BR. RATIO',7X,'NBRNCH',4X,'IBRNCH')
C      DO 15 I=1,NISO
C      DO 10 J=1,4
C      LANAB(I,J)=0.
C      10 CONTINUE
C      ISWTR=0
C      READ(50,1009) TAG(I),TR(I),ICLASS(I),LANAB(I,2),P(I),
C      1 NBRNCH(I),IBRNCH(I),ILAB
C      IF(ILAB.EQ.1) READ(5,1005) LANAB(I,1),LANAB(I,3),NIAB(I,4)
C      1005 FORMAT(3F10.0)
C
C      NAT 1180
C      NAT 1190
C      NAT 1200
C      NAT 1210
C      NAT 1220
C      NAT 1230
C      NAT 1231
C      NAT 1240
C      NAT 1250
C      NAT 1251
C      NAT 1260
C      NAT 1270
C      NAT 1280
C      NAT 1290
C      NAT 1300
C      NAT 1310
C      NAT 1320
C      NAT 1330
C      NAT 1340
C      NAT 1350
C      NAT 1360
C      NAT 1370
C      NAT 1380
C      NAT 1390
C      NAT 1400
C      NAT 1410
C      NAT 1420
C      NAT 1430
C      NAT 1440
C      NAT 1441
C      NAT 1450
C      NAT 1460
C      NAT 1470
C      NAT 1480
C      NAT 1490
C      NAT 1500
C      NAT 1510
C      NAT 1520
C      NAT 1530
C      NAT 1540
C      NAT 1550
C      NAT 1560
C      NAT 1570
C      NAT 1580
C      NAT 1590
C      NAT 1600
C      NAT 1601
C      NAT 1610
C      NAT 1620
C      NAT 1630
C      NAT 1640
C      NAT 1650
C      NAT 1650
C      NAT 1661
C      NAT 1670
C      NAT 1680

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      IF(P(I).EQ.0.) P(I)=1.
      WRITE(10,1006) TAG(I),TR(I),ICLASS(I),LAMB(I,2),P(I),
1  NBRNCH(I),IBRNCH(I)
1006 FORMAT(' ',A8,1PE12.3,4X,A4,2E10.3,2I10)
      IF (ILAB.EQ.1) WRITE(10,1007) LAMB(I,1),LAMB(I,3),LAMB(I,4)
1007 FORMAT(' ',21X,'LAMB-S',1PE10.3,/' ',19X,'LAMB-OLI',E10.3, /
1  ' ',19X,'LAMB-LLI',E10.3)
15  CONTINUE
1008 FORMAT(A8,2X,E10.0,8X,A2,2F10.0,2I2,2X,I2)
      WRITE(16) (TAG(I),I=1,NISO)
C
C  SET LAHR
C
      DO 20 I=1,NISO
      LAHR(I)=PLN/TR(I)
20  CONTINUE
C
C  CHECK CLASS; SET LAMX AND PRAC
C
      IF (IOPT.EQ.0) CALL LUNG1(ICLASS,X0,NX0)
C
C
C  CALCULATE ENTRIES OF COEFFICIENT MATRIX
C  ENTER NON-ZERO ENTRIES BY ROWS TO FORM VECTOR A
C
      IROW=0
      IKOUNT=0
      ICOL1=0
      ICOL=0
      DO 30 K=1,NISO
C  K - INDEX OF RADIONUCLIDE
C  J - INDEX OF COMPARTMENT
      J=0
      ICOL=ICOL+1
25  CONTINUE
C
C  LUNG COMPARTMENTS
C
      IF (IOPT.EQ.0) CALL LUNG2(A,RINDEX,CINDEX,IBOW,ICOL,ICOL1,
1  IKOUNT,LAHR,P,K,J,N,NBRNCH,IBRNCH)
C
C
C  GI TRACT
C
      IF (IOPT.EQ.0.OR.IOPT.EQ.1) CALL GI(A,RINDEX,CINDEX,IBOW,
1  ICOL,ICOL1,IKOUNT,LAHR,P,K,J,N,NBRNCH,IBRNCH,X0,NX0)
C
C
C  OTHER ORGANS
C
      IF (IOPT.EQ.0.OR.IOPT.EQ.1.OR.IOPT.EQ.2) CALL ORGAN(A,RINDEX,
1  CINDEX,I*OW,ICOL,ICOL1,IKOUNT,LAHR,P,K,J,N,NBRNCH,IBRNCH,
2  ISORS,X0,NX0)

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```

NAT 1690
NAT 1700
NAT 1701
NAT 1710
NAT 1720
NAT 1730
NAT 1731
NAT 1740
NAT 1750
NAT 1760
NAT 1770
NAT 1780
NAT 1790
NAT 1800
NAT 1810
NAT 1820
NAT 1830
NAT 1840
NAT 1850
NAT 1860
NAT 1870
NAT 1880
NAT 1890
NAT 1900
NAT 1910
NAT 1920
NAT 1930
NAT 1940
NAT 1950
NAT 1960
NAT 1970
NAT 1980
NAT 1990
NAT 2000
NAT 2010
NAT 2020
NAT 2030
NAT 2040
NAT 2050
NAT 2060
NAT 2061
NAT 2070
NAT 2080
NAT 2090
NAT 2100
NAT 2110
NAT 2120
NAT 2121
NAT 2130
NAT 2140
NAT 2150
NAT 2160
NAT 2170
NAT 2180
NAT 2181
NAT 2182

```

C	IF (K.EQ.1) GO TO 30	NAT 2190
	CALL INDEX (ICOL1, NCON, K, NBRNCH(K), IBBNCH(K))	NAT 2200
30	CONTINUE	NAT 2210
	IF (IOPT.EQ.0) NORGPB=NCRG+9	NAT 2220
	IF (IOPT.EQ.1) NORGPB=NORG+8	NAT 2230
	IF (IOPT.EQ.2) NORGPB=NCRG+9	NAT 2240
	IF (IOPT.EQ.0) IORG=1	NAT 2250
	IF (IOPT.EQ.1) IORG=1	NAT 2260
	IF (IOPT.EQ.2) IORG=1	NAT 2270
	WRITE (16) (ISOPS(I), I=IORG, NORGPB)	NAT 2280
C		NAT 2290
C	DETERMINE TOTAL NUMBER OF COMPARTMENTS (SUM1)	NAT 2300
C		NAT 2310
	SUM1=0.00	NAT 2320
	DO 35 K=1, NISO	NAT 2330
	SUM1=SUM1+NCON(K)	NAT 2340
35	CONTINUE	NAT 2350
C		NAT 2360
C		NAT 2370
C	PRINT COEFFICIENT MATRIX BY ROWS	NAT 2380
C		NAT 2390
	WRITE (51,1009)	NAT 2400
1009	FORMAT ('MATRIX A PRINTED BY ROWS')	NAT 2410
	II=0	NAT 2420
	DO 55 I=1, SUM1	NAT 2430
	III=0	NAT 2440
	DO 40 J=1, SUM1	NAT 2450
	PRNT(J)=0.00	NAT 2460
40	CONTINUE	NAT 2470
	WRITE (51,1010)	NAT 2480
1010	FORMAT ('0')	NAT 2490
45	CONTINUE	NAT 2500
	II=II+1	NAT 2510
	III=III+1	NAT 2520
	IF (RINDEX(II).EQ.I) GO TO 50	NAT 2530
	II=II-1	NAT 2540
	III=III-1	NAT 2550
	INDXA(2,2*I-1)=-1*III	NAT 2560
	INDXA(2,2*I)=8*III	NAT 2570
	IDIAG(I)=II	NAT 2580
	WRITE (51,1011) (PRNT(J), J=1, SUM1)	NAT 2590
1011	FORMAT (' ',8D15.6)	NAT 2600
	GO TO 55	NAT 2610
50	CONTINUE	NAT 2620
	PRNT(CINDEX(II))=A(II)	NAT 2630
	INDXA(1,II)=8*(CINDEX(II)-1)	NAT 2640
	GO TO 45	NAT 2650
55	CONTINUE	NAT 2660
C		NAT 2670
	DO 60 I=1, NISO	NAT 2680
	LB(I)=-LAWR(I)	NAT 2690
	NBLK(I)=NCON(I)	NAT 2700
60	CONTINUE	NAT 2710
	NOSUBN=NISO	NAT 2720
	N=SUM1	NAT 2730
		NAT 2740


```

      NPLA=I*DIAG(M)
C
C PRINT NON-ZERO ELEMENTS OF THE COEFFICIENT MATRIX AND
C THE ROW AND COLUMN IN WHICH EACH ONE APPEARS
C
      WRITE(6,1012)
1012 FORMAT('1',3X,'ENTRIES OF A',2X,'ROW',2X,'COL')
      DO 65 I=1,NPLA
      WRITE(51,1013) A(I),RINDEX(I),CINDEX(I)
1013 FORMAT(' ',D15.6,2I5)
65 CONTINUE
C
      RETURN
      END

```

```

SAT 2750
SAT 2760
SAT 2770
SAT 2780
SAT 2790
SAT 2800
SAT 2810
SAT 2820
SAT 2830
SAT 2840
SAT 2850
SAT 2860
SAT 2870
SAT 2880

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      SUBROUTINE LONG(D3,D4,D5)
C
C THIS SUBROUTINE DETERMINES THE ENTRIES OF THE COEFFICIENT
C MATRIX CORRESPONDING TO THE LONG COMPARTMENTS
C
C A(NT(10))
C   A(NT(I)) = LAMB(K)*PRAC(K,I) FOR PATHWAY I, DAUGHTER K
C
C D(9) - REGIONAL DEPOSITION VALUES
C   D(1) = D3      D(5) = D5
C   D(2) = D3      D(6) = D5
C   D(3) = D4      D(7) = D5
C   D(4) = D4      D(8) = D5
C
C P(9,3)
C   P(X,I) = REGIONAL FRACTION TO PATHWAY
C             X = A,B,C,...J
C             I = 1 CLASS D
C               2 CLASS W
C               3 CLASS Y
C
C PPAC(10,9)
C   PPAC(I,J) = REGIONAL FRACTION IN PATHWAY J OF DAUGHTER I
C
C IC(10)
C   IC(I) = CLASS INDEX FOR DAUGHTER I
C           1 - CLASS D
C           2 - CLASS W
C           3 - CLASS Y
C
C LAMBDA(3,9)
C   LAMBDA(J,X) = LW 2/(REMOVAL HALFLIFE FOR PATHWAY

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```

LONG 10
LONG 20
LONG 30
LONG 40
LONG 50
LONG 60
LONG 70
LONG 80
LONG 90
LONG 100
LONG 110
LONG 120
LONG 130
LONG 140
LONG 150
LONG 160
LONG 170
LONG 180
LONG 190
LONG 200
LONG 210
LONG 220
LONG 230
LONG 240
LONG 250
LONG 260
LONG 270
LONG 280
LONG 290
LONG 300
LONG 310
LONG 320

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15	CONTINUE	LONG 770
	IF (ICLASS(I).NE.YEAR) GO TO 20	LONG 780
	IC(I)=3	LONG 790
	GO TO 25	LONG 800
20	CONTINUE	LONG 810
	WRITE(10,1001)	LONG 820
1001	FORMAT('0',11X,' ERROR. CLASS DOES NOT COMPARE')	LONG 830
	CALL EPROR	LONG 840
25	CONTINUE	LONG 850
C		LONG 860
C	SET LAMB AND PRAC	LONG 870
C		LONG 880
	DO 30 J=1,9	LONG 890
	LAMB(I,J)=LAMBDA(IC(I),J)	LONG 900
	PRAC(I,J)=P(J,IC(I))	LONG 910
30	CONTINUE	LONG 920
	LAMB(I,10)=0. DO	LONG 930
C		LONG 940
C	IF PARENT IS CLASS Y AND DAUGHTER IS CLASS D OR W	LONG 950
C	SET LAMB(I,10) TO LAMB(I,9)	LONG 960
C		LONG 970
	IF(I.EQ.1) GO TO 35	LONG 980
	IF (IC(I-1).EQ.3. AND. (IC(I).EQ.1.OR.IC(2).EQ.2))	LONG 990
	1 LAMB(I,10)=LAMB(I,9)	LONG 991
35	CONTINUE	LONG 1000
	IF(I.GT.1) GO TO 45	LONG 1010
C		LONG 1020
C	SET INITIAL VALUES FOR ODE	LONG 1030
C		LONG 1040
	DO 40 K=1,NX0	LONG 1050
	XO(K)=0. DO	LONG 1060
	IF(K.LE.8) XO(K)=D(K)*PRAC(1,K)	LONG 1070
40	CONTINUE	LONG 1080
45	CONTINUE	LONG 1090
C		LONG 1100
	RETURN	LONG 1110
C		LONG 1120
	ENTRY LONG2(A,RINDEX,CINDEX,IROW,ICOL,ICD,1,	LONG 1130
	1 IKOUNT,LAMB,P,K,J,N,NBRCH,IBRCH)	LONG 1131
C		LONG 1140
C	LONG COMPARTMENTS	LONG 1150
C	PATHWAYS A-H	LONG 1160
C		LONG 1170
	IF(K.EQ.1) GO TO 55	LONG 1180
	DO 50 I=1,9	LONG 1190
	AMT(I)=LAMB(K)*PRAC(K,I)	LONG 1200
50	CONTINUE	LONG 1210
	AMT(10)=LAMB(K)*(1. DO-PRAC(K,9))	LONG 1220
55	CONTINUE	LONG 1230
	J=J+1	LONG 1240
	IROW=IBRW+1	LONG 1250
	IF(K.EQ.1) GO TO 70	LONG 1260
	Q=P(K)	LONG 1270
60	CONTINUE	LONG 1280
	IF(J.LE.2) N=2	LONG 1290
	IF(J.EQ.3.OR.J.EQ.4) N=2	LONG 1300

	IP(J.GE.5.AND.J.LE.9) N=8	LOW 1310
	DO 65 I=1,N	LOW 1320
	IKOUNT=IKOUNT+1	LOW 1330
	ICOL=ICOL+1	LOW 1340
	A(IKOUNT)=AMT(J)*P(K)	LOW 1350
	RINDEX(IKOUNT)=IROW	LOW 1360
	CINDEX(IKOUNT)=ICOL	LOW 1370
65	CONTINUE	LOW 1380
	IF(NBRNCH(K).EQ.IBRNCH(K).AND.NBRNCH(K).GT.1)	LOW 1390
	1 CALL CHANGE(P(K),K,ICOL,ICOL1,NCOR,J,N,IOPT,660)	LOW 1391
	P(K)=Q	LOW 1400
	II=2*N-J	LOW 1410
	IF(J.LE.2) II=N-J	LOW 1420
	ICOL=ICOL+NCOR(K-1)-II	LOW 1430
	IF(NBRNCH(K).EQ.3.AND.IBRNCH(K).EQ.2) ICOL=ICOL+NCOR(K-2)	LOW 1440
	GO TO 75	LOW 1450
70	CONTINUE	LOW 1460
	ICOL=ICOL+1	LOW 1470
75	CONTINUE	LOW 1480
	IKOUNT=IKOUNT+1	LOW 1490
	A(IKOUNT)=-LAX(K,J)	LOW 1500
	RINDEX(IKOUNT)=IROW	LOW 1510
	CINDEX(IKOUNT)=ICOL	LOW 1520
	IF(J.EQ.9.AND.K.GT.1) ICOL=ICOL+J/N*N	LOW 1530
	IF(J.EQ.9) GO TO 80	LOW 1540
	IF(K.EQ.1) GO TO 55	LOW 1550
	ICOL=ICOL+J/N*N	LOW 1560
	GO TO 55	LOW 1570
80	CONTINUE	LOW 1580
C		LOW 1590
C	PATHWAYS I AND J	LOW 1600
C		LOW 1610
	IROW=IROW+1	LOW 1620
	J=J+1	LOW 1630
	IF(K.EQ.1) GO TO 95	LOW 1640
	Q=P(K)	LOW 1650
85	CONTINUE	LOW 1660
	DO 90 I=1,2	LOW 1670
	ICOL=ICOL+1	LOW 1680
	IKOUNT=IKOUNT+1	LOW 1690
	A(IKOUNT)=AMT(J)*P(K)	LOW 1700
	RINDEX(IKOUNT)=IROW	LOW 1710
	CINDEX(IKOUNT)=ICOL	LOW 1720
90	CONTINUE	LOW 1730
	IF(NBRNCH(K).EQ.IBRNCH(K).AND.NBRNCH(K).GT.1)	LOW 1740
	1 CALL CHANGE(P(K),K,ICOL,ICOL1,NCOR,J,N,IOPT,635)	LOW 1741
	P(K)=Q	LOW 1750
	ICOL=ICOL+NCOR(K-1)-2	LOW 1760
	IF(NBRNCH(K).EQ.3.AND.IBRNCH(K).EQ.2) ICOL=ICOL+NCOR(K-2)	LOW 1770
	GO TO 100	LOW 1780
95	CONTINUE	LOW 1790
	ICOL=9	LOW 1800
100	CONTINUE	LOW 1810
	IKOUNT=IKOUNT+1	LOW 1820
	IF(J.EQ.9) A(IKOUNT)=LAX(K,8)*PRAC(K,9)	LOW 1830
	IF(J.EQ.10) A(IKOUNT)=LAX(K,8)*(1.-PRAC(K,9))	LOW 1840

	RIINDEX(IKOUNT)=IROW	LUN 1850
	CINDEX(IKOUNT)=ICOL	LUN 1860
	II=1	LUN 1870
	IF(J.EQ.10) II=2	LUN 1880
	ICOL=ICCL+II	LUN 1890
	IKOUNT=IKOUNT+1	LUN 1900
	IF(J.EQ.9) A(IKOUNT)=-LANX(K,J)	LUN 1910
	IF(J.EQ.10) A(IKOUNT)=0.00	LUN 1920
	RIINDEX(IKOUNT)=IROW	LUN 1930
	CINDEX(IKOUNT)=ICOL	LUN 1940
	IF(J.EQ.10.AND.K.GT.1) ICOL=ICOL+J/2*2	LUN 1950
	IF(J.EQ.10) GO TO 105	LUN 1960
	IF(K.EQ.1) GO TO 80	LUN 1970
	ICOL=ICOL+J/2*2	LUN 1980
	GO TO 80	LUN 1990
105	CONTINUE	LUN 2000
C		LUN 2010
C	PATHWAYS K AND L	LUN 2020
C		LUN 2030
	DO 125 I=1,2	LUN 2040
	IROW=IROW+1	LUN 2050
	J=J+1	LUN 2060
	IF(K.EQ.1) GO TO 115	LUN 2070
	Q=P(K)	LUN 2080
110	CONTINUE	LUN 2090
	ICOL=ICOL+1	LUN 2100
	IKOUNT=IKOUNT+1	LUN 2110
	A(IKOUNT)=LANX(K)+P(K)	LUN 2120
	RIINDEX(IKOUNT)=IROW	LUN 2130
	CINDEX(IKOUNT)=ICOL	LUN 2140
	IF(NBRCH(K).EQ.IBRCH(K).AND.NBRCH(K).GT.1)	LUN 2150
	1 CALL CHANGE(P(K),K,ICOL,ICOL1,NCOR,J,B,IPT,0110)	LUN 2151
	P(K)=Q	LUN 2160
	ICOL=ICOL+NCOR(K-1)-5	LUN 2170
	IF(NBRCH(K).EQ.3.AND.IBRCH(K).EQ.2) ICOL=ICOL+NCOR(X-2)	LUN 2180
	GO TO 120	LUN 2190
115	CONTINUE	LUN 2200
	ICOL=5+I	LUN 2210
120	CONTINUE	LUN 2220
	IKOUNT=IKOUNT+1	LUN 2230
	A(IKOUNT)=LANX(K,5+I)	LUN 2240
	RIINDEX(IKOUNT)=IROW	LUN 2250
	CINDEX(IKOUNT)=ICOL	LUN 2260
	IKOUNT=IKOUNT+1	LUN 2270
	ICOL=ICCL+5	LUN 2280
	A(IKOUNT)=-LANX(K,4)	LUN 2290
	RIINDEX(IKOUNT)=IROW	LUN 2300
	CINDEX(IKOUNT)=ICOL	LUN 2310
	IF(K.EQ.1) GO TO 125	LUN 2320
	ICOL=ICOL+J	LUN 2330
125	CONTINUE	LUN 2340
	RETURN	LUN 2350
	END	LUN 2360

```

SUBROUTINE GI (A, RINDEX, CINDEX, IROW, ICOL, ICOL1, IKOUNT, LAMB,
1 P, K, J, N, IBBRNC, IBBRNC, X0, NX0)
C
C THIS SUBROUTINE DETERMINES THE ENTRIES OF THE COEFFICIENT
C MATRIX CORRESPONDING TO THE COMPARTMENTS OF THE GI TRACT
C
C GILAN (4)
C GILAN (I) = LAMBDA FOR COMPARTMENT I OF GI TRACT
C
C LAMB (10,4)
C LAMB (I, 1) = LAMBDA OF ABSORPTION IN S FOR DAUGHTER I
C LAMB (I, 2) = LAMBDA OF ABSORPTION IN SI FOR DAUGHTER I
C LAMB (I, 3) = LAMBDA OF ABSORPTION IN OLI FOR DAUGHTER I
C LAMB (I, 4) = LAMBDA OF ABSORPTION IN LLI FOR DAUGHTER I
C
C IMPLICIT REAL*8 (A-H,O-Z)
C REAL*8 LAMB (10,10), LAMB (1), LAMB (10,4)
C INTEGER*2 RINDEX (1), CINDEX (1)
C DIMENSION A (1), GILAN (4), P (1), IBBRNC (1), IBBRNC (1), X0 (1)
C
C COMMON TABLES/ LAMB (10,4), LAMB (10,10), ORGIB (10), TAG (10),
1 ICON (10,10), IOPT, SCOR (10), NISO, NORG
C
C DATA GILAN/24.00, 6.00, 1.8861538860, 1.00/
C
C SET INITIAL VALUES FOR ODE
C
C IF (K.GT.1) GO TO 15
C IF (IOPT.EQ.0.OR.IOPT.EQ.2) GO TO 15
C DO 10 I=1, NX0
C X0 (I) = 0.00
C IF (I.EQ.1) X0 (I) = 1.00
10 CONTINUE
C
15 CONTINUE
C
C GI TRACT
C STOMACH
C
C IROW=IROW+1
C J=J+1
C IF (K.EQ.1) GO TO 25
C Q=P (K)
20 CONTINUE
C ICOL=ICOL+1
C IKOUNT=IKOUNT+1
C A (IKOUNT)=LAMB (K)*P (K)
C RINDEX (IKOUNT)=IROW
GI 10
GI 11
GI 20
GI 30
GI 40
GI 50
GI 60
GI 70
GI 80
GI 90
GI 100
GI 110
GI 120
GI 130
GI 140
GI 150
GI 160
GI 170
GI 180
GI 190
GI 200
GI 201
GI 210
GI 220
GI 230
GI 240
GI 250
GI 260
GI 270
GI 280
GI 290
GI 300
GI 310
GI 320
GI 330
GI 340
GI 350
GI 360
GI 370
GI 380
GI 390
GI 400
GI 410
GI 420
GI 430
GI 440
GI 450
GI 460
GI 470

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	CI INDEX (IKOUNT) = ICOL	GI 480
	IF (NBRNCH(K) .EQ. IBRNCH(K) .AND. NBRNCH(K) .GT. 1)	GI 490
	1 CALL CHANGE(P(K), K, ICOL, ICGL1, NCON, J, N, IOPT, 620)	GI 495
	P(K) = Q	GI 500
	IF (IOPT .EQ. 0) ICOL = ICOL + NCON(K-1) - 1	GI 510
	IF (IOPT .EQ. 1) ICOL = ICOL + NCON(K-1)	GI 520
	IF (NBRNCH(K) .EQ. 3 .AND. IBRNCH(K) .EQ. 2) ICOL = ICOL + NCON(K-2)	GI 530
	GO TO 30	GI 540
25	CONTINUE	GI 550
	IF (IOPT .EQ. 0) ICOL = 2	GI 560
	IF (IOPT .EQ. 1) ICOL = 1	GI 570
30	CONTINUE	GI 580
	IF (IOPT .EQ. 1) GO TO 45	GI 590
	DO 35 I = 2, 4, 2	GI 600
	IKOUNT = IKOUNT + 1	GI 610
	A(IKOUNT) = LANX(K, I)	GI 620
	RI INDEX (IKOUNT) = IROW	GI 630
	CI INDEX (IKOUNT) = ICOL	GI 640
	ICOL = ICOL + I	GI 650
35	CONTINUE	GI 660
	ICOL = ICOL + 3	GI 670
	DO 40 I = 1, 2	GI 680
	IKOUNT = IKOUNT + 1	GI 690
	A(IKOUNT) = LANX(K, 4)	GI 700
	RI INDEX (IKOUNT) = IROW	GI 710
	CI INDEX (IKOUNT) = ICOL	GI 720
	ICOL = ICOL + 1	GI 730
40	CONTINUE	GI 740
45	CONTINUE	GI 750
	IKOUNT = IKOUNT + 1	GI 760
	A(IKOUNT) = -GILAN(1)	GI 770
	A(IKOUNT) = A(IKOUNT) - LANAB(K, 1)	GI 780
	RI INDEX (IKOUNT) = IROW	GI 790
	CI INDEX (IKOUNT) = ICOL	GI 800
	IF (K .EQ. 1) GO TO 50	GI 810
	ICOL = ICOL + J	GI 820
50	CONTINUE	GI 830
C		GI 840
C	SI, OLI, LIT	GI 850
C		GI 860
	DO 70 I = 1, 3	GI 870
	IROW = IROW + 1	GI 880
	J = J + 1	GI 890
	IF (K .EQ. 1) GO TO 60	GI 900
	Q = P(K)	GI 910
55	CONTINUE	GI 920
	ICOL = ICOL + 1	GI 930
	IKOUNT = IKOUNT + 1	GI 940
	A(IKOUNT) = LANB(K)	GI 950
	RI INDEX (IKOUNT) = IROW	GI 960
	CI INDEX (IKOUNT) = ICOL	GI 970
	A(IKOUNT) = LANB(K) * P(K)	GI 980
	IF (NBRNCH(K) .EQ. IBRNCH(K) .AND. NBRNCH(K) .GT. 1)	GI 990
	1 CALL CHANGE(P(1), K, ICOL, ICOL1, NCON, J, N, IOPT, 655)	GI 995
	P(K) = Q	GI 1000
	ICOL = ICOL + NCON(K-1) - 1	GI 1010

	IF (NBRNCH(K) .EQ. 3.AND. IBRNCH(K) .EQ.2) ICOL=ICOL+BCCN(K-2)	GI 1020
	70 TO 65	GI 1030
60	CONTINUE	GI 1040
	ICOL=J-1	GI 1050
65	CONTINUE	GI 1060
	IKOUNT=IKOUNT+1	GI 1070
	A(IKOUNT)=GILAM(I)	GI 1080
	RINDEX(IKOUNT)=IROW	GI 1090
	CINDEX(IKOUNT)=ICOL	GI 1100
	ICOL=ICOL+1	GI 1110
	IKOUNT=IKOUNT+1	GI 1120
	A(IKOUNT)=-GILAM(I+1)	GI 1130
	A(IKOUNT)=A(IKOUNT)-LAMAB(K,I+1)	GI 1140
	RINDEX(IKOUNT)=IROW	GI 1150
	CINDEX(IKOUNT)=ICOL	GI 1160
	IF (K.EQ.1) GO TO 70	GI 1170
	ICOL=ICOL+J	GI 1180
70	CONTINUE	GI 1190
C		GI 1200
	RETURN	GI 1210
	END	GI 1220

	SUBROUTINE ORGAN (A,RINDEX,CINDEX,IROW,ICOL,ICOL1,IKOUNT,	ORGAN 10
	1 LAMR,P,K,J,N,NBRNCH,IBRNCH,ISORS,IO,IXO)	ORGAN 11
C		ORGAN 20
C	THIS SUBROUTINE DETERMINES THE ENTRIES OF THE COEFFICIENT	ORGAN 30
C	MATRIX CORRESPONDING TO THE COMPARTMENTS OF OTHER ORGANS OF	ORGAN 40
C	THE BODY	ORGAN 50
C		ORGAN 60
C		ORGAN 70
C	ICON(10,10)	ORGAN 80
C	ICON(I,J) = NO. OF COMPARTMENT FOR DAUGHTER I,	ORGAN 90
C	ORGAN J	ORGA 100
C	AS(10,10), TBS(10,10)	ORGA 110
C	AS(I,J) = COEF. OF EXPONENTIAL OF COMPARTMENT I, ORGAN J	ORGA 120
C	TBS(I,J) =POWER OF E, COMPARTMENT I ORGAN J	ORGA 130
C		ORGA 140
C	IS(10)	ORGA 150
C	IS(I) - FRACTION OF THE DAUGHTER RETURNING	ORGA 160
C	TO BLOOD FOR ORGAN I	ORGA 170
C		ORGA 180
	IMPLICIT REAL*8 (A-H,O-Z)	ORGA 190
	REAL*8 LAMX(10,10),LAMR(1),LAMAB(10,4)	ORGA 200
	INTEGER*2 RINDEX(1),CINDEX(1)	ORGA 210
	DIMENSION A(1),AS(10,10),TBS(10,10),IS(10),ISORS(1),P(1),	ORGA 220
	1 IBRNCH(1),IBRNCH(1),IO(1)	ORGA 221
C		ORGA 230

	COMMON/TABLES/ LANAB (10,4), LANI (10, 10), ORGHH (10), TAG (10),	ORGA 240
	1 ICON (10, 10), IOPT, NCON (10), NISO, NORG	ORGA 241
	COMMON/PAK21/ SGH	ORGA 250
C		ORGA 260
C		ORGA 270
C	ORDER ORGANS	ORGA 280
C		ORGA 290
	IO=0	ORGA 300
	IF (NORG.EQ.0) GO TO 80	ORGA 310
	IF (K.GT.1) READ (50,1001) (IS (I), I=1, NORG)	ORGA 320
1001	FORMAT (10I2)	ORGA 330
	WRITE (10,1002)	ORGA 340
1002	FORMAT ('0', //, ' ISO IOPE', 4X, 'ORGAN', 5X, 'A'S AND TB'S')	ORGA 350
	WRITE (10,1003) TAG (K)	ORGA 360
1003	FORMAT (' ', A3)	ORGA 370
	SNORN=0.00	ORGA 380
	DO 65 I=1, NORG	ORGA 390
	READ (50, 1004) ISORS (I+8), ORGHH (I), ICON (K, I)	ORGA 400
1004	FORMAT (I2, A8, I2)	ORGA 410
	ICONK I=ICON (K, I)	ORGA 420
	READ (50,1005) (AS (JJ, I), JJ=1, ICONKI), (TRS (JJ, I), JJ=1, ICONKI)	ORGA 430
1005	FORMAT (9F10.0)	ORGA 440
	WRITE (10, 1006) ORGHH (I), (AS (JJ, I), JJ=1, ICONKI),	ORGA 450
	1 (TRS (JJ, I), JJ=1, ICONKI)	ORGA 451
1006	FORMAT (' ', 10X, A8, 2X, 'PRE 12.3, /, (' ', 20X, 8E12.3)	ORGA 460
	SUM=0.00	ORGA 470
	DO 10 JJ=1, ICONKI	ORGA 480
	SUM=SUM+AS (JJ, I)	ORGA 490
10	CONTINUE	ORGA 500
	SNORN=SNORN+SUM	ORGA 510
	DO 60 HH=1, ICONKI	ORGA 520
	IROW=IROW+1	ORGA 530
	J=J+1	ORGA 540
	IF (K.EQ.1) GO TO 30	ORGA 550
	Q=P (K)	ORGA 560
	ISW=1	ORGA 570
15	CONTINUE	ORGA 580
	ISW=ISW+1	ORGA 590
	DO 25 JJ=1, NORG	ORGA 600
	ICONKH=ICON (K-1, JJ)	ORGA 610
	IF (IBRNCH (K).GT.1.AND.ISW.EQ.2) ICONKH=ICON (K-2, JJ)	ORGA 620
	DO 25 KH=1, ICONKH	ORGA 630
	CALL PAKB	ORGA 640
	ICOL=ICOL+1	ORGA 650
	IKOUNT=IKOUNT+1	ORGA 660
	A (IKOUNT)=IS (JJ) *LANR (K) *AS (HH, I) *P (K)	ORGA 670
	IF (IS (JJ).EQ.1) GO TO 20	ORGA 680
	IF (JJ.EQ.I) A (IKOUNT)=A (IKOUNT)+(1.00-IS (JJ;) *	ORGA 690
	1 LANR (K) *AS (HH, I) /SUM *P (K)	ORGA 691
20	CONTINUE	ORGA 700
	RINDEX (IKOUNT)=IROW	ORGA 710
	CINDEX (IKOUNT)=ICOL	ORGA 720
25	CONTINUE	ORGA 730
	IF (NBRNCH (K).EQ.IBRNCH (K).AND.NBRNCH (K).GT.1;	ORGA 740
	1 CALL CHANGE (P (K), K, ICOL, ICOL1, NCON, J, N, IOPT, 6 15)	ORGA 741
	P (K)=Q	ORGA 750

	IF (NB BRCH(K) .EQ. 3. AND. IBRCH(K) .EQ. 2) ICOL=ICOL+BCOR(K-1)	ORG 760
	GO TO 35	ORG 770
30	CONTINUE	ORG 780
	ICOL=0	ORG 790
35	CONTINUE	ORG 800
	ICOL=ICOL+1	ORG 810
	IF (IOPT.EQ.1) GO TO 45	ORG 820
	IF (IOPT.EQ.2) GO TO 55	ORG 830
	DO 40 JJ=1,5,2	ORG 840
	IKOUNT=IKOUNT+1	ORG 850
	A (IKOUNT)=LAMY(K, JJ)*AS(MH, I)	ORG 860
	RINDEX (IKOUNT)=IPOR	ORG 870
	CINDEX (IKOUNT)=ICOL	ORG 880
	ICOL=ICOL+2	ORG 890
40	CONTINUE	ORG 900
	ICOL=ICOL+2	ORG 910
	IKOUNT=IKOUNT+1	ORG 920
	A (IKOUNT)=LAMY(K, 9)*AS(MH, I)	ORG 930
	RINDEX (IKOUNT)=IKOR	ORG 940
	CINDEX (IKOUNT)=ICOL	ORG 950
	ICOL=ICOL+4	ORG 960
45	CONTINUE	ORG 970
	DO 50 JJ=1,4	ORG 980
	IKOUNT=IKOUNT+1	ORG 990
	A (IKOUNT)=LAMAB(K, JJ)*BS(MH, I)	ORG 1000
	RINDEX (IKOUNT)=IPOR	ORG 1010
	CINDEX (IKOUNT)=ICOL	ORG 1020
	ICOL=ICOL+1	ORG 1030
50	CONTINUE	ORG 1040
55	CONTINUE	ORG 1050
	IO=IO+1	ORG 1060
	ICOL=ICOL-1+IO	ORG 1070
	IKOUNT=IKOUNT+1	ORG 1080
	A (IKOUNT)=-TBS(MH, I)	ORG 1090
	RINDEX (IKOUNT)=IPOR	ORG 1100
	CINDEX (IKOUNT)=ICOL	ORG 1110
	IF (K.EQ.1) GO TO 60	ORG 1120
	IF (IOPT.EQ.0) ICOL=ICOL+16	ORG 1130
	IF (IOPT.EQ.1) ICOL=ICOL+4	ORG 1140
	IF (IOPT.EQ.2) ICOL=ICOL	ORG 1150
60	CONTINUE	ORG 1160
65	CONTINUE	ORG 1170
C		ORG 1180
C	SET INITIAL VALUES FOR ODE	ORG 1190
C		ORG 1200
	IF (K.GT.1) GO TO 80	ORG 1210
	IF (IOPT.EQ.0. OR. IOPT.EQ.1) GO TO 80	ORG 1220
	Y=0	ORG 1230
	DO 70 INO=1, NORG	ORG 1240
	ICORRI=ICOR(K, INO)	ORG 1250
	DO 70 ICH=1, ICORRI	ORG 1260
	I=I+1	ORG 1270
	YO (I)=AS (ICH, INO)/SBORN	ORG 1280
70	CONTINUE	ORG 1290
	IOP1=IO+1	ORG 1300
	DO 75 I=IOP1, NIO	ORG 1310

	IO(I) = 0.00	ORG 1320
75	CONTINUE	ORG 1330
80	CONTINUE	ORG 1340
	IF (IOPT.EQ.3) NCON(K) = IO+16	ORG 1350
	IF (IOPT.EQ.4) NCON(K) = IO+4	ORG 1360
	IF (IOPT.EQ.2) NCON(K) = IO	ORG 1370
	RETURN	ORG 1380
	END	ORG 1390

	SUBROUTINE YNA(Y,N)	YNA 10
	IMPLICIT REAL*8 (A-H,O-Z)	YNA 20
C		YNA 30
C	THIS SUBROUTINE PRINTS MICROCURIE-DAYS FOR EACH	YNA 40
C	TIME INTERVAL	YNA 50
C		YNA 60
C	NORG - NUMBER OF ORGANS (EXCLUDING LUNG AND GI)	YNA 70
C	ICON(I,J) - NO. OF COMPARTMENTS FOR DAUGHTER I, ORGAN J	YNA 80
C	NCON(I) - NO. OF COMPARTMENTS FOR DAUGHTER I	YNA 90
C	NISO - NUMBER OF ELEMENTS IN THE CHAIN	YNA 100
C	LANX(I,J) - LW2/(HALFTIME OF DAUGHTER I IN COMPARTMENT J)	YNA 110
C	ORGN(I) - NAME OF ORGAN I	YNA 120
C	TAG(I) - ATOMIC SYMBOL AND ATOMIC NUMBER FOR	YNA 130
C	DAUGHTER I	YNA 140
C		YNA 150
	REAL*8 TAG,LANX,X(10,21),ORGN,LANAB	YNA 160
	COMMON/TABLES/ LANAB(10,4), LANX(10,10),ORGN(10),TAG(10),	YNA 170
	I ICON(10,10), IOPT,NCON(10), NISO,NORG	YNA 171
	COMMON/PRINT/TPRL(10),DTPR(10),TPRINT	YNA 180
	DI MENSION Y(1)	YNA 190
C		YNA 200
C	PRINT SOLUTION VECTOR Y	YNA 210
C		YNA 220
	PRINT 1001,TPRINT,(Y(I),I=1,N)	YNA 230
	1001 FORMAT('0T='1PE16.9,3X'Y FOLLOWS'/(1P6R19.10))	YNA 240
	K=0	YNA 250
C		YNA 260
C	X(I,1) - BP REGION FOR DAUGHTER I	YNA 270
C	X(I,2) - TB REGION FOR DAUGHTER I	YNA 280
C	X(I,3) - P REGION FOR DAUGHTER I	YNA 290
C	X(I,4) - LYMPH FOR DAUGHTER I	YNA 300
C	X(I,5) - STOMACH FOR DAUGHTER I	YNA 310
C	X(I,6) - SI FOR DAUGHTER I	YNA 320
C	X(I,7) - ULI FOR DAUGHTER I	YNA 330
C	X(I,8) - LLI FOR DAUGHTER I	YNA 340
C	X(I,J) - COMPARTMENT J (J=9,10,...18) FOR DAUGHTER I	YNA 350
C	X(I,19) - MICROCURIES ENTERING BLOOD FROM LUNGS FOR DAUGHTER I	YNA 360
C	X(I,20) - MICROCURIES ENTERING GI FROM LUNGS FOR DAUGHTER I	YNA 370

C	X(I,2) - MICROCURIES ENTERING BLOOD FROM GI FOR DAUGHTER I	YNA	390
C		YNA	390
	DO 45 I=1,NISO	YNA	400
	IF (IOPT.EQ.1.OR.IOPT.EQ.2) GO TO 10	YNA	410
	X(I,1)=Y(1+K)+I(2+K)	YNA	420
	X(I,2)=Y(3+K)+Y(4+K)+Y(11+K)+Y(12+K)	YNA	430
	X(I,3)=Y(5+K)+Y(6+K)+Y(7+K)+Y(8+K)	YNA	440
	X(I,4)=Y(9+K)+Y(10+K)	YNA	450
	GO TO 15	YNA	460
10	CONTINUE	YNA	470
	X(I,1)=0.	YNA	480
	X(I,2)=0.	YNA	490
	X(I,3)=0.	YNA	500
	X(I,4)=0.	YNA	510
15	CONTINUE	YNA	520
	IF (IOPT.EQ.0) IORG=12	YNA	530
	IF (IOPT.EQ.1) IORG=0	YNA	540
	IF (IOPT.EQ.2) GO TO 25	YNA	550
	DO 20 J=1,4	YNA	560
	X(I,J+4)=Y(IORG+K+J)	YNA	570
20	CONTINUE	YNA	580
	GO TO 10	YNA	590
25	CONTINUE	YNA	600
	X(I,5)=0.	YNA	610
	X(I,6)=0.	YNA	620
	X(I,7)=0.	YNA	630
	X(I,8)=0.	YNA	640
30	CONTINUE	YNA	650
	IF (IORG.EQ.0) GO TO 40	YNA	660
	ICONIJ=0	YNA	670
	DO 35 J=1,NIORG	YNA	680
	II=ICONIJ+1	YNA	690
	ICONIJ=ICOH(I,J)+ICONIJ	YNA	700
	X(I,J+8)=0.DO	YNA	710
	IF (IOPT.EQ.0) IORG=16	YNA	720
	IF (IOPT.EQ.1) IORG=0	YNA	730
	IF (IOPT.EQ.2) IORG=0	YNA	740
	DO 35 M=II,ICONIJ	YNA	750
	X(I,J+8)=X(I,J+8)+Y(IORG+K+M)	YNA	760
35	CONTINUE	YNA	770
40	CONTINUE	YNA	780
	IF (IOPT.EQ.0) X(I,19)=Y(1+K)*LANX(I,1)+Y(3+K)*LANX(I,3)	YNA	790
	+Y(5+K)*LANX(I,5)+Y(9+K)*LANX(I,9)	YNA	791
	IF (IOPT.EQ.0) X(I,20)=Y(2+K)*LANX(I,2)+Y(4+K)*LANX(I,4)	YNA	800
	+Y(11+K)*LANX(I,4)+Y(12+K)*LANX(I,4)	YNA	801
	IF (IOPT.EQ.0.OR.IOPT.EQ.1) X(I,21)=X(I,5)*LANAB(I,1)	YNA	810
	+X(I,6)*LANAB(I,2)+X(I,7)*LANAB(I,3)+X(I,8)*LANAB(I,4)	YNA	811
	K=K+NIORG(I)	YNA	820
45	CONTINUE	YNA	830
	WRITE(10,1002) TPRINT	YNA	840
1002	FORMAT('IT = ',1PE10.4)	YNA	850
	IF (IOPT.EQ.0) WRITE(10,1003) TAG(1)	YNA	860
1003	FORMAT('OM: MICROCURIES-DAYS FROM INHALATION OF ',A8)	YNA	870
	IF (IOPT.EQ.1) WRITE(10,1004) TAG(1)	YNA	880
1004	FORMAT('OM: MICROCURIE-DAYS FROM INGESTION OF ',A8)	YNA	890
	IF (IOPT.EQ.2) WRITE(10,1005) TAG(1)	YNA	900

1005	FORMAT('0MICROCURIE-DAYS FROM INJECTION OF ',A8)	YBA	910
	IF(IOPT.EQ.0) WRITE(10,1006)	YBA	920
1006	FORMAT('0LUNGS',59I,'MICROCURIES ENTERING'/	YBA	930
1	' ',67I,'BLOOD',7I,'GI'/' ISOTOPE',6I,'B-P',9I,'T-B',10I,	YBA	931
2	'P',9I,'LYMPH')	YBA	932
	DO 50 I=1,NISO	YBA	940
	IF(IOPT.EQ.0) WRITE(10,1007) TAG(I), (X(I,J),J=1,4),X(I,19),X(I,20)	YBA	950
50	CONTINUE	YBA	960
1007	FORMAT(' ',A3,' (2X,1PE10.4), 10I,E10.4, 2X,E10.4	YBA	970
	IF(IOPT.EQ.0.OR.IOPT.EQ.1) WRITE(10,1008)	YBA	980
1008	FORMAT('0'/'0GI TRACT',56I,'MICROCURIES ENTERING'/	YBA	990
1	'BLOOD'/' ISOTOPE',4I,'STOMACH',7I,'S.I.',7I,'U.L.I.',6I,	YBA	991
2	'L.I.I.')	YBA	992
	DO 55 I=1,NISO	YBA	1000
	IF(IOPT.EQ.0.OR.IOPT.EQ.1) WRITE(10,1007) TAG(I),	YBA	1010
1	(X(I,J),J=5,8),X(I,21)	YBA	1011
55	CONTINUE	YBA	1020
	NORGP8=NORG+9	YBA	1030
	IF(NORG.EQ.0) GO TO 65	YBA	1040
	WRITE(10,1009) (ORGN(I),I=1,NORG)	YBA	1050
1009	FORMAT('0'/'0OTHER ORGANS'/'0 ISOTOPE',5I,10(A8,4I))	YBA	1060
	DO 60 I=1,NISO	YBA	1070
	WRITE(10,1010) TAG(I), (X(I,J),J=9,NORGP8)	YBA	1080
1010	FORMAT(' ',A8,10(1X,1PE11.4))	YBA	1090
60	CONTINUE	YBA	1100
	CALL OTHER(X)	YBA	1110
65	CONTINUE	YBA	1120
	WRITE(16) IPRINT	YBA	1130
	IF(IOPT.EQ.0) IORG=1	YBA	1140
	IF(IOPT.EQ.1) IORG=1	YBA	1150
	IF(IOPT.EQ.2) IORG=1	YBA	1160
	DO 70 I=1,NISO	YBA	1170
	WRITE(16) (X(I,J),J=IORG,NORGP8)	YBA	1180
	IF(IOPT.EQ.0) WRITE(16) X(I,19),X(I,20),X(I,21)	YBA	1190
	WRITE(7,1011) (X(I,J),J=IORG,NORGP8)	YBA	1200
1011	FORMAT(1PE10.3)	YBA	1210
70	CONTINUE	YBA	1220
	RETURN	YBA	1230
	END	YBA	1240

	SUBROUTINE OTHER (X)	OTHER	10
C		OTHER	20
C	PERFORMS OTHER TISSUES TRANSFORMATION	OTHER	30
C		OTHER	40
	REAL*8 TAG,LANI,X(10,1),ORGN,TBODY/T. BODY '/,LANAB	OTHER	50
	COMMON/TABLES/ LANAB(10,4),LANI(10,10),ORGN(10),TAG(10),	OTHER	60
	1 ICON('0,10),IOPT,ICON(10),NISO,NORG	OTHER	61

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REAL THASS(25)/200.,250.,400.,200.,135.,310.,1800.,85.,0.,
1 1000.,0.,29000.,11.,100.,4000.,1000.,1500.,1500.,2600.,180.,35.,
2 20.,70000.,0.,14./
DIMENSION IOTHER(10),OTHER(10)
NORGH1=NORG-1
READ(50,1001) MOTOP,(IOther(I),I=1,NORGH1)
1001 FORMAT(40I2)
IF(MOTOP.EQ.0) RETURN
SHASS=0.
DO 10 I=1,NORGH1
SHASS=SHASS+THASS(IOther(I))
10 CONTINUE
DO 15 J=1,NISO
OTHER(J)=X(I,R+NORG)
X(I,R+NORG)=OTHER(J)/(THASS(23)-SHASS)*THASS(23)
DO 15 J=1,NORGH1
X(I,R+J)=X(I,R+J)-THASS(IOther(J))*OTHER(J)/
1 (THASS(23)-SHASS)
15 CONTINUE
WRITE(10,1002)
1002 FORMAT('CHAPTER TRANSFORMATION')
WRITE(10,1003) (ORGN(I),I=1,NORGH1),TBD
1003 FORMAT(' ',12X,10(A8,4X))
NORGP8=NORG+8
DO 20 I=1,NISO
WRITE(10,1004) TAG(I),(X(I,J),J=9,NORGP8)
1004 FORMAT(' ',A8,10(1X,1P811.8))
20 CONTINUE
RETURN
END

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OTHER 70
OTHER 71
OTHER 72
OTHER 80
OTHER 90
OTHER 160
OTHER 110
OTHER 120
OTHER 130
OTHER 140
OTHER 150
OTHER 160
OTHER 170
OTHER 180
OTHER 190
OTHER 200
OTHER 210
OTHER 211
OTHER 220
OTHER 230
OTHER 240
OTHER 250
OTHER 260
OTHER 270
OTHER 280
OTHER 290
OTHER 300
OTHER 310
OTHER 320
OTHER 330

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SUBROUTINE INDEX(ICOL1,NCON,K,IN,II)
C
C DETERMINES FIRST COLUMN FOR THE DAUGHTER
C
DIMENSION NCON(1)
IF(IN.GE.2) GO TO 10
ICOL1=ICOL1+NCON(K-1)
RETURN
10 CONTINUE
IF(IN.EQ.11) GO TO 15
IF(II.EQ.1) RETURN
ICOL1=ICOL1+NCON(K-2)
RETURN
15 CONTINUE
ICOL1=ICOL1+NCON(K-1)+NCON(K-2)
RETURN

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INDEX 10
INDEX 20
INDEX 30
INDEX 40
INDEX 50
INDEX 60
INDEX 70
INDEX 80
INDEX 90
INDEX 100
INDEX 110
INDEX 120
INDEX 130
INDEX 140
INDEX 150
INDEX 160


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AXEB  TITLE 'SOLVE AX=B WHERE A IS A COMPACTED LOWER TRI. MATRIX'  AXEB  10
AXEB  START 0  AXEB  20
* CALL AXEB(A,X,B,INDXA,N)  AXEB  30
* A IS AN N BY N LOWER TRIANGULAR MATRIX. THE VECTOR A CONSISTS  AXEB  40
* OF THE NONZERO ENTRIES IN A STORED SEQUENTIALLY BY ROWS.  AXEB  50
* SOLVE LINEAR SYSTEM OF EQUATIONS.  AXEB  60
***** IT IS ASSUMED THAT THE MAX. NO. OF NONZERO ENTRIES IN A ROW OF A  AXEB  70
***** IS .LE. 52.  HOWEVER, THE DIAG. ENTRIES ARE NONZERO.  AXEB  80
***** AT MOST 51 SUBRANS IN SUBL  AXEB  90
A      EQU  1  AXEB 100
X      EQU  2  AXEB 110
B      EQU  3  AXEB 120
INDXA  EQU  4  AXEB 130
INDIA  EQU  5  AXEB 140
IINC   EQU  6  AXEB 150
I      EQU  7  AXEB 160
J      EQU  8  AXEB 170
BADDR  EQU  9  AXEB 180
        EQU 10  AXEB 190
        USING AXEB,15  AXEB 190
        STB  14,12,12(13)  AXEB 200
        L4  1,5,0(1)  AXEB 210
        LA  IINC, 4  AXEB 220
        L  7,0(5)  B  AXEB 230
        LA  0,16  AXEB 240
        SLA 7,3  8W  AXEB 250
        SR  7,IINC  8(N-1)  AXEB 260
        SR  I,I  AXEB 270
        LR  5,4  AXEB 280
        LA  BADDR,POINT+14  AXEB 290
        SP  A,IINC  A=ADD(A(K-1,K-1),K=1,....,B  AXEB 300
LOOP   SDR  0,0  FORN SUBL=SUB (J=1,....,I-1) A(I,J)*K(J)  AXEB 310
        FOR I=1,....,N.  AXEB 320
        L  11,4(INDIA)  AXEB 330
        B  0(BADDR,11)  AXEB 340
        ..  AXEB 350
        ..  AXEB 360
        L  J,400(INDXA)  AT LEAST 51  AXEB 370
        LD  6,408(A)  AXEB 380
        RD  6,0(X,J)  AXEB 390
        SDR  0,6  AXEB 400
        L  J,392(INDXA)  AT LEAST 50  AXEB 410
        LD  4,400(A)  AXEB 420
        RD  4,0(X,J)  AXEB 430
        SDR  0,4  AXEB 440
        L  J,384(INDXA)  AT LEAST 49  AXEB 450
        LD  2,392(A)  AXEB 460
        RD  2,0(X,J)  AXEB 470

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SDR	0,2			AKEB	480
L	J,376 (INDIA)	AT LEAST	48	AKEB	490
LD	6,384 (A)			AKEB	500
ND	6,0 (X,J)			AKEB	510
SDR	0,6			AKEB	520
L	J,368 (INDIA)	AT LEAST	47	AKEB	530
LD	4,376 (A)			AKEB	540
ND	4,0 (X,J)			AKEB	550
SDR	0,4			AKEB	560
L	J,360 (INDIA)	AT LEAST	46	AKEB	570
LD	2,368 (A)			AKEB	580
ND	2,0 (X,J)			AKEB	590
SDR	0,2			AKEB	600
L	J,352 (INDIA)	AT LEAST	45	AKEB	610
LD	6,360 (A)			AKEB	620
ND	4,0 (X,J)			AKEB	630
SDR	0,6			AKEB	640
L	J,344 (INDIA)	AT LEAST	44	AKEB	650
LD	4,352 (A)			AKEB	660
ND	4,0 (X,J)			AKEB	670
SDR	0,4			AKEB	680
L	J,336 (INDIA)	AT LEAST	43	AKEB	690
LD	2,344 (A)			AKEB	700
ND	2,0 (X,J)			AKEB	710
SDR	0,2			AKEB	720
L	J,328 (INDIA)	AT LEAST	42	AKEB	730
LD	6,336 (A)			AKEB	740
ND	6,0 (X,J)			AKEB	750
SDR	0,6			AKEB	760
L	J,320 (INDIA)	AT LEAST	41	AKEB	770
LD	4,328 (A)			AKEB	780
ND	4,0 (X,J)			AKEB	790
SDR	0,4			AKEB	800
L	J,312 (INDIA)	AT LEAST	40	AKEB	810
LD	2,320 (A)			AKEB	820
ND	2,0 (X,J)			AKEB	830
SDR	0,2			AKEB	840
L	J,304 (INDIA)	AT LEAST	39	AKEB	850
LD	6,312 (A)			AKEB	860
ND	6,0 (X,J)			AKEB	870
SDR	0,6			AKEB	880
L	J,296 (INDIA)	AT LEAST	38	AKEB	890
LD	4,304 (A)			AKEB	900
ND	4,0 (X,J)			AKEB	910
SDR	0,4			AKEB	920
L	J,288 (INDIA)	AT LEAST	37	AKEB	930
LD	2,296 (A)			AKEB	940
ND	2,0 (X,J)			AKEB	950
SDR	0,2			AKEB	960
L	J,280 (INDIA)	AT LEAST	36	AKEB	970
LD	6,288 (A)			AKEB	980
ND	6,0 (X,J)			AKEB	990
SDR	0,6			AKEB	1000
L	J,272 (INDIA)	AT LEAST	35	AKEB	1010
LD	4,280 (A)			AKEB	1020
ND	4,0 (X,J)			AKEB	1030

SDR	0, 4			AXE	1040
L	J, 254 (INDXA)	AT LEAST	30	AXE	1050
LD	2, 272 (A)			AXE	1060
MD	2, 0 (X, J)			AXE	1070
SDR	0, 2			AXE	1080
L	J, 256 (INDXA)	AT LEAST	31	AXE	1090
LD	6, 264 (A)			AXE	1100
MD	6, 0 (X, J)			AXE	1110
SDR	0, 6			AXE	1120
L	J, 208 (INDXA)	AT LEAST	32	AXE	1130
LD	4, 256 (A)			AXE	1140
MD	4, 0 (X, J)			AXE	1150
SDR	0, 4			AXE	1160
L	J, 240 (INDXA)	AT LEAST	31	AXE	1170
LD	2, 248 (A)			AXE	1180
MD	2, 0 (X, J)			AXE	1190
SDR	0, 2			AXE	1200
L	J, 232 (INDXA)	AT LEAST	30	AXE	1210
LD	6, 240 (A)			AXE	1220
MD	6, 0 (X, J)			AXE	1230
SDR	0, 6			AXE	1240
L	J, 224 (INDXA)	AT LEAST	29	AXE	1250
LD	4, 232 (A)			AXE	1260
MD	4, 0 (X, J)			AXE	1270
SDR	0, 4			AXE	1280
L	J, 216 (INDXA)	AT LEAST	28	AXE	1290
LD	2, 224 (A)			AXE	1300
MD	2, 0 (X, J)			AXE	1310
SDR	0, 2			AXE	1320
L	J, 208 (INDXA)	AT LEAST	27	AXE	1330
LD	6, 216 (A)			AXE	1340
MD	6, 0 (X, J)			AXE	1350
SDR	0, 6			AXE	1360
L	J, 200 (INDXA)	AT LEAST	26	AXE	1370
LD	4, 208 (A)			AXE	1380
MD	4, 0 (X, J)			AXE	1390
SDR	0, 4			AXE	1400
L	J, 192 (INDXA)	AT LEAST	25	AXE	1410
LD	2, 200 (A)			AXE	1420
MD	2, 0 (X, J)			AXE	1430
SDR	0, 2			AXE	1440
L	J, 184 (INDXA)	AT LEAST	24	AXE	1450
LD	6, 192 (A)			AXE	1460
MD	6, 0 (X, J)			AXE	1470
SDR	0, 6			AXE	1480
L	J, 176 (INDXA)	AT LEAST	23	AXE	1490
LD	4, 184 (A)			AXE	1500
MD	4, 0 (X, J)			AXE	1510
SDR	0, 4			AXE	1520
L	J, 168 (INDXA)	AT LEAST	22	AXE	1530
LD	2, 176 (A)			AXE	1540
MD	2, 0 (X, J)			AXE	1550
SDR	0, 2			AXE	1560
L	J, 160 (INDXA)	AT LEAST	21	AXE	1570
LD	6, 168 (A)			AXE	1580
MD	6, 0 (X, J)			AXE	1590

SDR	0,6			AKR 1600
L	J, 152 (INDXA)	AT LEAST 20		AKR 1610
LD	4, 160 (A)			AKR 1620
ND	4, 0 (X, J)			AKR 1630
SDR	0,4			AKR 1640
L	J, 144 (INDXA)	AT LEAST 19		AKR 1650
LD	2, 152 (A)			AKR 1660
ND	2, 0 (X, J)			AKR 1670
SDR	0,2			AKR 1680
L	J, 136 (INDXA)	AT LEAST 18		AKR 1690
LD	6, 144 (A)			AKR 1700
ND	6, 0 (X, J)			AKR 1710
SDR	0,6			AKR 1720
L	J, 128 (INDXA)	AT LEAST 17 NONZERO OFF-DIAG. ENTRIES		AKR 1730
LD	4, 136 (A)			AKR 1740
ND	4, 0 (X, J)			AKR 1750
SDR	0,4			AKR 1760
L	J, 120 (INDXA)	AT LEAST 16 NONZERO OFF-DIAG. ENTRIES		AKR 1770
LD	2, 128 (A)			AKR 1780
ND	2, 0 (X, J)			AKR 1790
SDR	0,2			AKR 1800
L	J, 112 (INDXA)	AT LEAST 15 NONZERO OFF-DIAG. ENTRIES		AKR 1810
LD	6, 120 (A)			AKR 1820
ND	6, 0 (X, J)			AKR 1830
SDR	0,6			AKR 1840
L	J, 104 (INDXA)	AT LEAST 14 NONZERO OFF-DIAG. ENTRIES		AKR 1850
LD	4, 112 (A)			AKR 1860
ND	4, 0 (X, J)			AKR 1870
SDR	0,4			AKR 1880
L	J, 096 (INDXA)	AT LEAST 13 NONZERO OFF-DIAG. ENTRIES		AKR 1890
LD	2, 104 (A)			AKR 1900
ND	2, 0 (X, J)			AKR 1910
SDR	0,2			AKR 1920
L	J, 088 (INDXA)	AT LEAST 12 NONZERO OFF-DIAG. ENTRIES		AKR 1930
LD	6, 096 (A)			AKR 1940
ND	6, 0 (X, J)			AKR 1950
SDR	0,6			AKR 1960
L	J, 080 (INDXA)	AT LEAST 11 NONZERO OFF-DIAG. ENTRIES		AKR 1970
LD	4, 088 (A)			AKR 1980
ND	4, 0 (X, J)			AKR 1990
SDR	0,4			AKR 2000
L	J, 072 (INDXA)	AT LEAST 10 NONZERO OFF-DIAG. ENTRIES		AKR 2010
LD	2, 080 (A)			AKR 2020
ND	2, 0 (X, J)			AKR 2030
SDR	0,2			AKR 2040
L	J, 64 (INDXA)			AKR 2050
LD	6, 72 (A)			AKR 2060
ND	6, 0 (X, J)			AKR 2070
SDR	0,6			AKR 2080
L	J, 56 (INDXA)			AKR 2090
LD	4, 64 (A)			AKR 2100
ND	4, 0 (X, J)			AKR 2110
SDR	0,4			AKR 2120
L	J, 48 (INDXA)			AKR 2130
LD	2, 56 (A)			AKR 2140
ND	2, 0 (X, J)			AKR 2150

SDR	0,2		AXE 216C
L	J,4) (INDXA)		AXE 2170
LD	6,49(A)		AXE 2180
MD	6,0 (X,J)		AXE 2190
SDR	0,6		AXE 2200
L	J,3) (INDXA)		AXE 2210
LD	4,40(A)		AXE 2220
MD	4,0 (X,J)		AXE 2230
SDR	0,4		AXE 2240
L	J,24 (INDXA)		AXE 2250
LD	2,32(A)		AXE 2260
MD	2,0 (X,J)		AXE 2270
SDR	0,2		AXE 2280
L	J,15 (INDXA)		AXE 2290
LD	6,28(A)		AXE 2300
MD	6,0 (X,J)		AXE 2310
SDR	0,6		AXE 2320
L	J,9 (INDXA)		AXE 2330
LD	4,16(A)		AXE 2340
MD	4,0 (X,J)		AXE 2350
SDR	0,4		AXE 2360
L	J,0 (INDXA)		AXE 2370
LD	2,9(A)		AXE 2380
MD	2,0 (X,J)		AXE 2390
SDR	0,2		AXE 2400
**			AXE 2410
POINT	L 11,12(IND14)	8*(NO. NONZERO ENTRIES IN ROW I OF A)	AXE 2420
	A9 A,11		AXE 2430
	AP INDXA,11		AXE 2440
	AR IND14,0		AXE 2450
	AD 0,0(B,I)	COULD CHECK FOR LOSS OF LEADING FIGURES.	AXE 2460
	DD 0,0(A)	A(I,I)	AXE 2470
	STD 0,0(X,I)		AXE 2480
	DXLE I,II NC,LOOP		AXE 2490
	LT 14,12,12(13)		AXE 2500
	B? 14		AXE 2510
	END		AXE 2520

YPAY	TITLE 'FORM YDOT = A*Y + X0'	YPAY 10
YPAY	START 0	YPAY 20
	* CALL YPAY (YDOT,A,Y,X0,INDXA,N)	YPAY 30
	* FORM YDOT=A*Y+X0	YPAY 40
	* WHERE A IS AN P BY N MATRIX. HOWEVER, ONLY THE NONZERO ENTRIES IN A	YPAY 50
	* ARE STORED IN THE VECTOR, A. STORAGE IN A IS BY ROWS	YPAY 60
	* VIEW INDXA AS AN INTEGER** VECTOR.	YPA : 70
	* THE NONZERO ENTRY A(I) BELONGS IN COLUMN (INDXA(2I-1)+8)/8	YPAY 80
	* INDXA(4I-2) = -14*(NO. NONZERO ENTRIES IN ROW I OF MATRIX A).	YPAY 90

* INDXA(4I) = 3*(. **)-
 * Y AND X0 ARE FULL VECTORS WITH N ELEMENTS
 * A ZERO POW IN A IS OKAY.

 * SET FOR AT MOST 52 NONZERO ENTRIES IN A ROW OF THE MATRIX A.
 ** IT IS ASSUMED THAT THE MAXIMUM NO. OF NONZERO ENTRIES IN A ROW OF
 ** A IS .LE. 52.

***** PROBABLY A REGULAR LOOP WOULD BE BETTER.

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YDOT EQU 1
A EQU 2
Y EQU 3
X0 EQU 4
INDXA EQU 5
IND14 EQU 6
I EQU 7
IINC EQU 8
J EQU 9
    USING YPAY, 15
    STM 14,12,12(13)
    LM 1,6,0(1)
    LA IINC,8
    SR I,I
    L 9,0(6)
    SLA 9,3
    LA 00,16
    SR 9,8
    LR 6,5
    LA 11,X0ADD
LOOP LD 0,0(X0,I)
    I 12,4(IND14)
    B 0(11,12)

```

I=0
 W
 8W
 8(N-1)
 USE 6 TO REFERENCE (4I) AND (4I-2)
 FORM YDOT(I) IN 0. =X0(I)
 -14*(NO. NONZERO EIR. IN ROW I OF A)

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L 1,408(INDXA) AT LEAST 52
LD 2,408(A)
MD 2,0(J,Y)
ADR 0,2
L J,400(INDXA) AT LEAST 51
LD 6,400(A)
MD 6,0(J,Y)
ADR 0,6
L J,392(INDXA) AT LEAST 50
LD 4,392(A)
MD 4,0(J,Y)
ADR 0,4
L J,384(INDXA) AT LEAST 49
LD 2,384(A)
MD 2,0(J,Y)
ADR 0,2
L J,376(INDXA) AT LEAST 48
LD 6,376(A)
MD 6,0(J,Y)
ADR 0,6
L J,368(INDXA) AT LEAST 47
LD 4,368(A)

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- YPAY 100
- YPAY 110
- YPAY 120
- YPAY 130
- YPAY 140
- YPAY 150
- YPAY 160
- YPAY 170
- YPAY 180
- YPAY 190
- YPAY 200
- YPAY 210
- YPAY 220
- YPAY 230
- YPAY 240
- YPAY 250
- YPAY 260
- YPAY 270
- YPAY 280
- YPAY 290
- YPAY 300
- YPAY 310
- YPAY 320
- YPAY 330
- YPAY 340
- YPAY 350
- YPAY 360
- YPAY 370
- YPAY 380
- YPAY 390
- YPAY 400
- YPAY 410
- YPAY 420
- YPAY 430
- YPAY 440
- YPAY 450
- YPAY 460
- YPAY 470
- YPAY 480
- YPAY 490
- YPAY 500
- YPAY 510
- YPAY 520
- YPAY 530
- YPAY 540
- YPAY 550
- YPAY 560
- YPAY 570
- YPAY 580
- YPAY 590
- YPAY 600
- YPAY 610
- YPAY 620
- YPAY 630
- YPAY 640
- YPAY 650

ND	4,0(J,Y)		YPAY 660
ADR	0,4		YPA Y 670
L	J,360(INDXA)	AT LEAST 46	YPAY 680
LD	2,360(A)		YPA Y 690
ND	2,0(J,Y)		YPAY 700
ADR	0,2		YPA Y 710
L	J,352(INDXA)	AT LEAST 45	YPAY 720
LD	6,352(A)		YPA Y 730
ND	6,0(J,Y)		YPAY 740
ADR	0,6		YPA Y 750
L	J,344(INDXA)	AT LEAST 44	YPAY 760
LD	4,344(A)		YPA Y 770
ND	4,0(J,Y)		YPAY 780
ADR	0,4		YPA Y 790
L	J,336(INDXA)	AT LEAST 43	YPAY 800
LD	2,336(A)		YPA Y 810
ND	2,0(J,Y)		YPAY 820
ADR	0,2		YPA Y 830
L	J,328(INDXA)	AT LEAST 42	YPAY 840
LD	6,328(A)		YPA Y 850
ND	6,0(J,Y)		YPAY 860
ADR	0,6		YPA Y 870
L	J,320(INDXA)	AT LEAST 41	YPAY 880
LD	4,320(A)		YPA Y 890
ND	4,0(J,Y)		YPAY 900
ADR	0,4		YPA Y 910
L	J,312(INDXA)	AT LEAST 40	YPAY 920
LD	2,312(A)		YPA Y 930
ND	2,0(J,Y)		YPAY 940
ADR	0,2		YPA Y 950
L	J,304(INDXA)	AT LEAST 39	YPAY 960
LD	6,304(A)		YPA Y 970
ND	6,0(J,Y)		YPAY 980
ADR	0,6		YPA Y 990
L	J,296(INDXA)	AT LEAST 38	YPA 1000
LD	4,296(A)		YPA 1010
ND	4,0(J,Y)		YPA 1020
ADR	0,4		YPA 1030
L	J,288(INDXA)	AT LEAST 37	YPA 1040
LD	2,288(A)		YPA 1050
ND	2,0(J,Y)		YPA 1060
ADR	0,2		YPA 1070
L	J,280(INDXA)	AT LEAST 36	YPA 1080
LD	6,280(A)		YPA 1090
ND	6,0(J,Y)		YPA 1100
ADR	0,6		YPA 1110
L	J,272(INDXA)	AT LEAST 35	YPA 1120
LD	4,272(A)		YPA 1130
ND	4,0(J,Y)		YPA 1140
ADR	0,4		YPA 1150
L	J,264(INDXA)	AT LEAST 34	YPA 1160
LD	2,264(A)		YPA 1170
ND	2,0(J,Y)		YPA 1180
ADR	0,2		YPA 1190
L	J,256(INDXA)	AT LEAST 33	YPA 1200
LD	6,256(A)		YPA 1210

MD	6,0 (J,Y)		YPA 1220
ADR	0,6		YPA 1230
L	J,248 (INDXA)	AT LEAST 32	YPA 1240
LD	4,248(A)		YPA 1250
MD	4,0 (J,Y)		YPA 1260
ADR	0,4		YPA 1270
L	J,240 (INDXA)	AT LEAST 31	YPA 1280
LD	2,240(A)		YPA 1290
MD	2,0 (J,Y)		YPA 1300
ADR	0,2		YPA 1310
L	J,232 (INDXA)	AT LEAST 30	YPA 1320
LD	6,232(A)		YPA 1330
MD	6,0 (J,Y)		YPA 1340
ADR	0,4		YPA 1350
L	J,224 (INDXA)	AT LEAST 29	YPA 1360
LD	4,224(A)		YPA 1370
MD	4,0 (J,Y)		YPA 1380
ADR	0,4		YPA 1390
L	J,216 (INDXA)	AT LEAST 28	YPA 1400
LD	2,216(A)		YPA 1410
MD	2,0 (J,Y)		YPA 1420
ADR	0,2		YPA 1430
L	J,208 (INDXA)	AT LEAST 27	YPA 1440
LD	6,208(A)		YPA 1450
MD	6,0 (J,Y)		YPA 1460
ADR	0,6		YPA 1470
L	J,200 (INDXA)	AT LEAST 26	YPA 1480
LD	4,200(A)		YPA 1490
MD	4,0 (J,Y)		YPA 1500
ADR	0,4		YPA 1510
L	J,192 (INDXA)	AT LEAST 25	YPA 1520
LD	2,192(A)		YPA 1530
MD	2,0 (J,Y)		YPA 1540
ADR	0,2		YPA 1550
L	J,184 (INDXA)	AT LEAST 24	YPA 1560
LD	6,184(A)		YPA 1570
MD	6,0 (J,Y)		YPA 1580
ADR	0,6		YPA 1590
L	J,176 (INDXA)	AT LEAST 23	YPA 1600
LD	4,176(A)		YPA 1610
MD	4,0 (J,Y)		YPA 1620
ADR	0,4		YPA 1630
L	J,168 (INDXA)	AT LEAST 22	YPA 1640
LD	2,168(A)		YPA 1650
MD	2,0 (J,Y)		YPA 1660
ADR	0,2		YPA 1670
L	J,160 (INDXA)	AT LEAST 21	YPA 1680
LD	6,160(A)		YPA 1690
MD	6,0 (J,Y)		YPA 1700
ADR	0,6		YPA 1710
L	J,152 (INDXA)	AT LEAST 20	YPA 1720
LD	4,152(A)		YPA 1730
MD	4,0 (J,Y)		YPA 1740
ADR	0,4		YPA 1750
L	J,144 (INDXA)	AT LEAST 19	YPA 1760
LD	2,144(A)		YPA 1770

ND	2,0(J,Y)		YPA 1780
ADR	0,2		YPA 1790
L	J,136(INDXA)	AT LEAST 18	YPA 1800
LD	6,136(A)		YPA 1810
ND	6,0(Y,J)		YPA 1820
ADR	0,6		YPA 1830
L	J,128(INDXA)	AT LEAST 17	YPA 1840
LD	4,128(A)		YPA 1850
ND	4,0(Y,J)		YPA 1860
ADR	0,4		YPA 1870
L	J,120(INDXA)	AT LEAST 16	YPA 1880
LD	2,120(A)		YPA 1890
ND	2,0(Y,J)		YPA 1900
ADR	0,2		YPA 1910
L	J,112(INDXA)	AT LEAST 15	YPA 1920
LD	6,112(A)		YPA 1930
ND	6,0(Y,J)		YPA 1940
ADR	0,6		YPA 1950
L	J,104(INDXA)	AT LEAST 14	YPA 1960
LD	4,104(A)		YPA 1970
ND	4,0(Y,J)		YPA 1980
ADR	0,4		YPA 1990
L	J,96(INDXA)	AT LEAST 13	YPA 2000
LD	2,96(A)		YPA 2010
ND	2,0(Y,J)		YPA 2020
ADR	0,2		YPA 2030
L	J,88(INDXA)	AT LEAST 12	YPA 2040
LD	6,88(A)		YPA 2050
ND	6,0(Y,J)		YPA 2060
ADR	0,6		YPA 2070
L	J,80(INDXA)	AT LEAST 11	YPA 2080
LD	4,80(A)		YPA 2090
ND	4,0(Y,J)		YPA 2100
ADR	0,4		YPA 2110
L	J,72(INDXA)	AT LEAST 10 NONZERO ENTRIES IN ROW I	YPA 2120
LD	2,72(A)		YPA 2130
ND	2,0(Y,J)		YPA 2140
ADR	0,2		YPA 2150
L	J,64(INDXA)	9	YPA 2160
LD	6,64(A)		YPA 2170
ND	6,0(Y,J)		YPA 2180
ADR	0,6		YPA 2190
L	J,56(INDXA)	8	YPA 2200
LD	4,56(A)		YPA 2210
ND	4,0(Y,J)		YPA 2220
ADR	0,4		YPA 2230
L	J,48(INDXA)	7	YPA 2240
LD	2,48(A)		YPA 2250
ND	2,0(Y,J)		YPA 2260
ADR	0,2		YPA 2270
L	J,40(INDXA)		YPA 2280
LD	6,40(A)		YPA 2290
ND	6,0(Y,J)		YPA 2300
ADR	0,6		YPA 2310
L	J,32(INDXA)	5	YPA 2320
LD	4,32(A)		YPA 2330

ND	4,0 (Y,J)		YPA 2340
ADR	0,4		YPA 2350
L	J,24 (INDXA)	4	YPA 2360
LD	2,24(A)		YPA 2370
ND	2,0 (Y,J)		YPA 2380
ADR	0,2		YPA 2390
L	J,16 (INDXA)	3	YPA 2400
LD	6,16(A)		YPA 2410
ND	6,0 (Y,J)		YPA 2420
ADR	0,6		YPA 2430
L	J,8 (INDXA)	2	YPA 2440
LD	4,8(A)		YPA 2450
ND	4,0 (Y,J)		YPA 2460
ADR	0,4		YPA 2470
L	J,00 (INDXA)		YPA 2480
LD	2,00(A)		YPA 2490
ND	2,00 (Y,J)		YPA 2500
ADR	0,2		YPA 2510
			YPA 2520
			YPA 2530
			YPA 2540
			YPA 2550
			YPA 2560
			YPA 2570
			YPA 2580
			YPA 2590
			YPA 2600
			YPA 2610

		AT LEAST 1 NONZERO ENTRY IN ROW I	
		8*(NO. NONZERO ENTRIES IN ROW I OF A)	

IOADD	L	12,12(IND14)	
	AR	A,12	
	AR	INDXA,12	
	AR	IND14,0	
	STD	0,0(YDOT,I)	
	BXIP	I,II MC,LOOP	
	LB	14,12,12(13)	
	BR	14	
	END		

INC002I	STOP	0
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APPENDIX B
SAMPLE INPUT AND OUTPUT

Included in this appendix are three examples of input and the corresponding output produced by the code. For each example we give the retention model, the input cards, and the output. The full printed output (unit 6 and unit 10) is given only for the first and third examples. Printout from unit 10 only is given for the second example.

Example 1

Radionuclide - CS-134

Half life - 2.05 years or 749. days

Mode of intake - inhalation (class D); see Fig. 2.1.

$$f_1 = 0.95$$

Source organs

1. Lungs
2. GIT

It is assumed that the total body retention is given by:

$$3. \text{ Total body: } R(t) = 0.13e^{-\frac{.693}{1.4}t} + 0.87e^{-\frac{.693}{135}t}$$

Print cumulated activity ($\mu\text{Ci-days}$) at these times:

2 da	5 yrs
7 da	10 yrs
30 da	20 yrs
60 da	30 yrs
180 da	40 yrs
1 yr	50 yrs

Below is an explanation of how Example 1 should be coded into card input. Follow Table 5.1, the description of the input data, and Table B.1, the data sheet, to fully understand the instructions.

Information Concerning the Differential Equations

Card 1 **NØ:** The number of differential equations.

12	Lungs
4	GIT
2	Total body
<hr/>	
18	

TLAST: The final value of T is 50 years or 18262.5 days.

All times must be expressed in days.

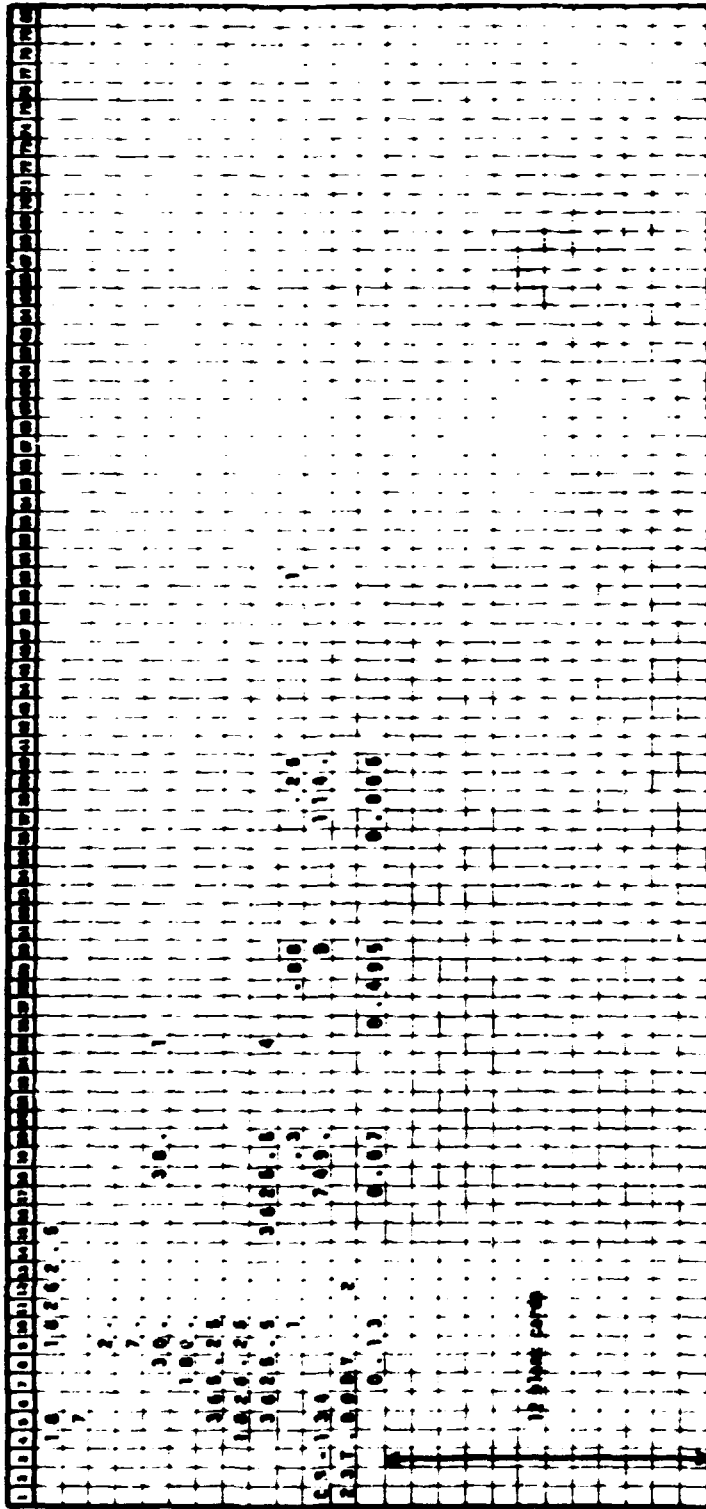
H1 and EPS are set to zero (left blank) therefore taking default values.

Card 2 **NPRT:** We wish to print cumulated activity ($\mu\text{Ci-days}$) at 12 different times; however, only 10 print intervals are allowable. Since some time periods are equally spaced, the time intervals may be considered as

<u>2.</u>	
<u>7.</u>	
<u>30.</u>	+ I·(30.) , where I=0,1
<u>180.</u>	
<u>365.25</u>	
<u>1826.5</u>	
<u>3652.5</u>	+ I·(3652.5) , where I=0,1,2,3,4

Thus there are 7 print intervals and TPRL(I), DTPR(I), NARPRT(I) are set equal to the underlined numbers as follows:

Table 8.1. CS-134: Incubate by Inhalation



	TPRL(I)	DTPR(I)	NARPRT(I)
<u>Card 3</u>	2.	0.	0
	7.	0.	0
.	30.	30.	1
.	180.	0.	0
.	365.25	0.	0
	1826.5	0.	0
<u>Card 9</u>	3652.5	3652.5	4

General Information for Chain of Radionuclides

Card 1 NISØ: 1 radionuclide in the chain
D3: .3)
D4: .08) ———— deposition fractions
D5: .25)
NØRG: 1 organ (excluding lungs and GIT)
IØPT: 0-inhalation
NØRAL: not applicable, set to 0

Card 2 TAG(1): CS-134
TR(1): 749. days
ICLASS(1): D
LAMAB(1,2): $6 \cdot 0.95 / (1 - 0.95) = 114.$
P(1),NBRNCH(1),IBRNCH(1): not applicable, set to 0
ILAB: 0

Card 3 omitted because ILAB=0

Description of Retention

Card 1 omitted for first nuclide in chain

Card 2 ISØRS(1): 23
ØRGNM(1): T.BØDY
ICØM(1,1): 2

Card 3 AS(1,1): 0.13
AS(2,1): 0.87
TBS(1,1): $0.693/1.4=0.495$
TBS(2,1): $0.693/135.=0.005$

Other Tissues Transformation

Since no transformation is to be performed, 1 blank card must be present for each time the cumulated activities ($\mu\text{Ci-days}$) are printed. Note this is 12, not 7.

Printout for Example 1 includes the following:

1. Summary of input concerning the differential equations.
2. Coefficient matrix printed by rows.
3. Nonzero elements of the coefficient matrix and the rows and columns where these elements appear.
4. The progress of the solution of the differential equations which includes the values of Y for each T.
5. Summary printout of input concerning the radionuclides and retention information for total body.
6. The remainder of the printout consists of cumulated activity for each organ at each time period. Printout for time periods 20 yrs., 30 yrs., 40 yrs., and 50 yrs. is omitted since the cumulated activities for these time periods are the same as those for 10 yrs. (3652.5 days).

Items 1-4 are output on unit 6; items 5 and 6 are output on unit 10.

OUTPUT FOR SAMPLE 1
CS-14 INTAKE BY SIMULATION

NO. POINT INTERVALS OBSERVED = 7 NO. READ = 7
POINTS SELECTED TO PRINT INTERVAL = 7, 10, 13 WHERE QMC, V,, N

Interval	QMC	V	...	N
2.4464000000 07	0.0			1
7.1400000000 08	0.0			3
8.0640000000 09	3.0000000000 09			1
1.9300000000 12	0.0			0
3.6524000000 12	0.0			0
1.9262500000 13	0.0			3
3.4527000000 13	3.4527000000 13			0

END 14
END 13
END 12
START 1.9262500000 13
END 3.4527000000 13
END 1.0000000000

ENTRIES OF A	NOB	CPL
-2.6030670 C2	1	1
-0.6030670 C2	2	2
-0.6030670 C2	3	3
-0.3005700 C1	4	4
-0.3005700 C1	5	5
0.0	6	6
0.0	7	7
-0.3005700 C1	8	8
-0.3005700 C1	9	9
-0.3005700 C1	10	10
0.0	11	11
0.0	12	12
-0.3005700 C1	13	13
-0.3005700 C1	14	14
-0.3005700 C1	15	15
-0.3005700 C1	16	16
-0.3005700 C1	17	17
-0.3005700 C1	18	18
-0.3005700 C1	19	19
-0.3005700 C1	20	20
0.0	21	21
0.0	22	22
-0.3005700 C1	23	23
-0.3005700 C1	24	24
-0.3005700 C1	25	25
-0.3005700 C1	26	26
-0.3005700 C1	27	27
-0.3005700 C1	28	28
-0.3005700 C1	29	29
-0.3005700 C1	30	30
0.0	31	31
0.0	32	32
-0.3005700 C1	33	33
-0.3005700 C1	34	34
-0.3005700 C1	35	35
-0.3005700 C1	36	36
-0.3005700 C1	37	37
-0.3005700 C1	38	38
-0.3005700 C1	39	39
-0.3005700 C1	40	40
0.0	41	41
0.0	42	42
-0.3005700 C1	43	43
-0.3005700 C1	44	44
-0.3005700 C1	45	45
-0.3005700 C1	46	46
-0.3005700 C1	47	47
-0.3005700 C1	48	48
-0.3005700 C1	49	49
-0.3005700 C1	50	50

KTAS = -1 FROM STEEP AT T = 3.0
 FROM TEST PATTERN WITH ABS(M) = 0.010

IT HAS BEEN DECIDED TO 0.000000000000000000-00 AND STEP WILL BE REPEATED

T= 2.00000000 C1	Y FOLLOWS	2.16001366930-03	1.09603359200-03	1.15272200510-03	1.35179077000-01	0.0
2.16001366930-03	2.16001366930-03	1.09603359200-03	1.15272200510-03	1.35179077000-01	0.0	0.0
0.0	2.16001366930-03	1.09603359200-03	1.15272200510-03	1.35179077000-01	0.0	0.0
0.01629132170-01	1.20320036000-03	0.16006700930-03	7.68051739600-03	1.63071561100-01	9.13359702330-01	0.0
T= 7.00000000 C1	Y FOLLOWS	2.16001366930-03	1.09603359200-03	1.15272200510-03	1.35179077000-01	0.0
2.16001366930-03	2.16001366930-03	1.09603359200-03	1.15272200510-03	1.35179077000-01	0.0	0.0
0.0	2.16001366930-03	1.09603359200-03	1.15272200510-03	1.35179077000-01	0.0	0.0
0.01629132170-01	1.20320036000-03	0.16006700930-03	7.68051739600-03	1.63071561100-01	9.13359702330-01	0.0
T= 3.00000000 C1	Y FOLLOWS	2.16001366930-03	1.09603359200-03	1.15272200510-03	1.35179077000-01	0.0
2.16001366930-03	2.16001366930-03	1.09603359200-03	1.15272200510-03	1.35179077000-01	0.0	0.0
0.0	2.16001366930-03	1.09603359200-03	1.15272200510-03	1.35179077000-01	0.0	0.0
0.01629132170-01	1.20320036000-03	0.16006700930-03	7.68051739600-03	1.63071561100-01	9.13359702330-01	0.0
T= 6.00000000 C1	Y FOLLOWS	2.16001366930-03	1.09603359200-03	1.15272200510-03	1.35179077000-01	0.0
2.16001366930-03	2.16001366930-03	1.09603359200-03	1.15272200510-03	1.35179077000-01	0.0	0.0
0.0	2.16001366930-03	1.09603359200-03	1.15272200510-03	1.35179077000-01	0.0	0.0
0.01629132170-01	1.20320036000-03	0.16006700930-03	7.68051739600-03	1.63071561100-01	9.13359702330-01	0.0
T= 1.00000000 C2	Y FOLLOWS	2.16001366930-03	1.09603359200-03	1.15272200510-03	1.35179077000-01	0.0
2.16001366930-03	2.16001366930-03	1.09603359200-03	1.15272200510-03	1.35179077000-01	0.0	0.0
0.0	2.16001366930-03	1.09603359200-03	1.15272200510-03	1.35179077000-01	0.0	0.0
0.01629132170-01	1.20320036000-03	0.16006700930-03	7.68051739600-03	1.63071561100-01	9.13359702330-01	0.0
T= 3.45250000 C2	Y FOLLOWS	2.16001366930-03	1.09603359200-03	1.15272200510-03	1.35179077000-01	0.0
2.16001366930-03	2.16001366930-03	1.09603359200-03	1.15272200510-03	1.35179077000-01	0.0	0.0
0.0	2.16001366930-03	1.09603359200-03	1.15272200510-03	1.35179077000-01	0.0	0.0
0.01629132170-03	1.20320036000-03	0.16006700930-03	7.68051739600-03	1.63071561100-01	9.13359702330-01	0.0
T= 1.02025000 C1	Y FOLLOWS	2.16001366930-03	1.09603359200-03	1.15272200510-03	1.35179077000-01	0.0
2.16001366930-03	2.16001366930-03	1.09603359200-03	1.15272200510-03	1.35179077000-01	0.0	0.0
0.0	2.16001366930-03	1.09603359200-03	1.15272200510-03	1.35179077000-01	0.0	0.0
0.01629132170-01	1.20320036000-03	0.16006700930-03	7.68051739600-03	1.63071561100-01	9.13359702330-01	0.0
T= 3.65200000 C3	Y FOLLOWS	2.16001366930-03	1.09603359200-03	1.15272200510-03	1.35179077000-01	0.0
2.16001366930-03	2.16001366930-03	1.09603359200-03	1.15272200510-03	1.35179077000-01	0.0	0.0
0.0	2.16001366930-03	1.09603359200-03	1.15272200510-03	1.35179077000-01	0.0	0.0
0.01629132170-01	1.20320036000-03	0.16006700930-03	7.68051739600-03	1.63071561100-01	9.13359702330-01	0.0

T= 7.30500C 10CD 03	Y FOLLOWS					
2.160013669 10-03	2.160013669 10-03	1.096033592 00-03	1.153007929 10-03	1.001732602 00-01	0.0	
C=J	7.600331506 00-02	3.601927015 00-02	0.0	0.0	3.0	
0.01620132170-03	1.203200360 00-03	0.160067609 30-03	7.600517396 00-03	1.030715611 60-01	9.133701700 10 01	
T= 1.095100000 00	Y FOLLOWS					
2.160013669 10-03	2.160013669 10-03	1.096033592 00-03	1.153007929 10-03	1.001732602 00-01	0.0	
C=J	7.600331506 00-02	3.601927015 00-02	0.0	0.0	3.0	
0.01620132170-03	1.203200360 00-03	0.160067609 30-03	7.600517396 00-03	1.030715611 60-01	9.133701602 10 01	
T= 1.001000000 00	Y FOLLOWS					
2.160013669 10-03	2.160013669 10-03	1.096033592 00-03	1.153007929 10-03	1.001732602 00-01	0.0	
C=J	7.600331506 00-02	3.601927015 00-02	0.0	0.0	3.0	
0.01620132170-03	1.203200360 00-03	0.160067609 30-03	7.600517396 00-03	1.030715611 60-01	9.133701605 10 01	
T= 1.025200000 00	Y FOLLOWS					
2.160013669 10-03	2.160013669 10-03	1.096033592 00-03	1.153007929 10-03	1.001732602 00-01	0.0	
C=J	7.600331506 00-02	3.601927015 00-02	0.0	0.0	3.0	
0.01620132170-03	1.203200360 00-03	0.160067609 30-03	7.600517396 00-03	1.030715611 60-01	9.133701605 90 01	

RPLAC = C

PERCENT DEPOSITION						
	H-P	T-B		P		
	0.30	0.00		0.25		
ISOTOPE	HAIFLIFE	CLASS	LABOR	NO. DAYS	DOSE	INDOSE
CS-134	7.6905 02	D	1.0000 02	1.0000 00	0	0

ISOTOPE	CYCLE	A'S	ART	TB'S
CS-134	2.9000	1.3000-01	0.7000-01	0.9500-01
				5.0000-01

T = 7.0000 00

MICROCUBIES-DAYS FROM INHALATION OF CS-134

LUNGS

ISOTOPE	H-P	T-B	P	LUNGS
CS-134	4.12400-03	2.24920-03	1.44970-01	2.75300-02

MICROCUBIES ENTERING
BLOOD

GI

4.51570-01 1.53990-01

GI TRACT

ISOTOPE	STOMACH	S.I.	U.L.I.	L.L.I.
CS-134	4.41470-03	1.29320-03	4.04030-03	5.50630-03

MICROCUBIES ENTERING
BLOOD

1.46290-01

OTHER ORGANS

ISOTOPE	T.BODY
CS-134	4.43200-01

T = 7.0000 00

MICROCUBIES-DAYS FROM INHALATION OF CS-134

LUNGS

ISOTOPE	H-P	T-B	P	LUNGS
CS-134	4.17000-03	2.25030-03	1.44210-01	3.59940-02

MICROCUBIES ENTERING
BLOOD

GI

4.75770-01 1.54000-01

GI TRACT

ISOTOPE	STOMACH	S.I.	U.L.I.	L.L.I.
CS-134	4.41470-03	1.29320-03	4.16050-03	7.67210-03

MICROCUBIES ENTERING
BLOOD

1.46290-01

OTHER ORGANS

ISOTOPE	T.BODY
CS-134	3.47000 00

OTHER ORGANS
ISOTOPE 1 BODY
CS-130 4.4995D 01

GI TRACT
ISOTOPE STOMACH
CS-130 4.163D-03 1.2812D-03 0.1665D-03 7.0005D-03
L.L.I. 0.L.I. L.L.I.

TISSUES
ISOTOPE B-P
CS-130 0.1200D-03 2.2503D-03 1.0022D-01 3.0019D-02
L L L L
MICROCURIES-DAYS FROM IRRADIATION OF CS-130
I = 1.0000D 02
MICROCURIES ENTERING
BLOOD GI
CS-130 0.7500D-01 1.5000D-01

OTHER ORGANS
ISOTOPE 1 BODY
CS-130 2.7190D 01

GI TRACT
ISOTOPE STOMACH
CS-130 0.163D-01 1.2912D-03 0.1635D-03 7.0005D-03
L.L.I. 0.L.I. L.L.I.

TISSUES
ISOTOPE B-P
CS-130 0.1200D-03 2.2503D-03 1.0022D-01 3.0019D-02
L L L L
MICROCURIES-DAYS FROM IRRADIATION OF CS-130
I = 6.3400D 01
MICROCURIES ENTERING
BLOOD GI
CS-130 0.7500D-01 1.5000D-01

OTHER ORGANS
ISOTOPE 1 BODY
CS-130 1.4700D 01

GI TRACT
ISOTOPE STOMACH
CS-130 0.163D-01 1.2912D-03 0.1635D-03 7.0005D-03
L.L.I. 0.L.I. L.L.I.

TISSUES
ISOTOPE B-P
CS-130 0.1200D-03 2.2503D-03 1.0022D-01 3.0019D-02
L L L L
MICROCURIES-DAYS FROM IRRADIATION OF CS-130
I = 3.0000D 01
MICROCURIES ENTERING
BLOOD GI
CS-130 0.7500D-01 1.5000D-01

T = 3.65250 02

MICROCURIES-DAYS FROM INHALATION OF CS-134

LUNGS

ISOTOPE	H-P	T-B	P	LYMPH
CS-134	4.32900-03	2.25030-03	1.80220-03	3.60190-02

MICROCURIES ENTERING
BLOOD GI

4.75800-01 1.54000-01

GI TRACT

ISOTOPE	STOMACH	S.I.	U.L.I.	L.L.I.
CS-134	4.41630-03	1.28320-03	4.16650-03	7.68650-03

MICROCURIES ENTERING
BLOOD

1.46290-01

OTHER ORGANS

ISOTOPE	T.BODY
CS-134	3.04990 01

T = 1.32630 03

MICROCURIES-DAYS FROM INHALATION OF CS-134

LUNGS

ISOTOPE	H-P	T-B	P	LYMPH
CS-134	4.32900-03	2.25030-03	1.80220-03	3.60190-02

MICROCURIES ENTERING
BLOOD GI

4.75800-01 1.54000-01

GI TRACT

ISOTOPE	STOMACH	S.I.	U.L.I.	L.L.I.
CS-134	4.41630-03	1.28320-03	4.16650-03	7.68650-03

MICROCURIES ENTERING
BLOOD

1.46290-01

OTHER ORGANS

ISOTOPE	T.BODY
CS-134	3.16990 01

T = 3.65250 01

MICROCURIES-DAYS FROM INHALATION OF CS-134

LUNGS

ISOTOPE	H-P	T-B	P	LYMPH
CS-134	4.32900-03	2.25030-03	1.80220-03	3.60190-02

MICROCURIES ENTERING
BLOOD GI

4.75900-01 1.54000-01

GI TRACT

ISOTOPE	STOMACH	S.I.	U.L.I.	L.L.I.
CS-134	4.41630-03	1.28320-03	4.16650-03	7.68650-03

MICROCURIES ENTERING
BLOOD

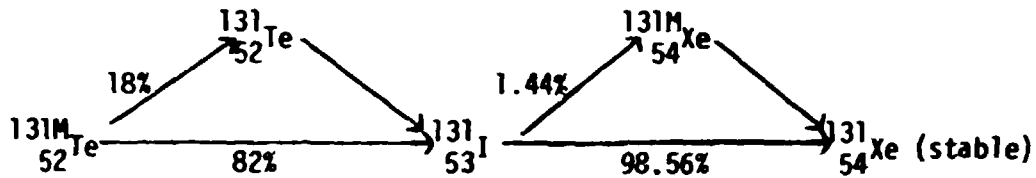
1.46290-01

OTHER ORGANS

ISOTOPE	T.BODY
CS-134	4.14030 01

Example 2

Chain of Radionuclides -



Half lives -	Te-131M	2. hours
	Te-131	25 minutes
	I-131	8.06 days
	Xe-131M	11.8 days

Mode of intake - inhalation (class D - all radionuclides)

Tellurium as a parent of Iodine:

$$f_1 = .25$$

Source organs

1. Lungs

2. GIT

$$3. \text{ Cort bone: } R(t) = 0.0183e^{-\frac{.693}{9.2}t} + 0.01e^{-\frac{.693}{8000}t}$$

$$4. \text{ Tra bone: } R(t) = 4.57 \times 10^{-3}e^{-\frac{.693}{9.2}t} + 0.01e^{-\frac{.693}{8000}t}$$

$$5. \text{ Kidney: } R(t) = 1.42 \times 10^{-3}e^{-\frac{.693}{9.2}t} + 0.034e^{-\frac{.693}{23}t}$$

$$6. \text{ Muscle: } R(t) = 0.128e^{-\frac{.693}{9.2}t} + 0.1e^{-\frac{.693}{17.7}t}$$

$$7. \text{ Liver: } R(t) = 8.23 \times 10^{-3}e^{-\frac{.693}{9.2}t} + 0.055e^{-\frac{.693}{10.2}t}$$

$$8. \text{ Thyroid: } R(t) = 9.14 \times 10^{-5}e^{-\frac{.693}{9.2}t}$$

9. Other tissue: $R(t) = 0.159e^{-\frac{.693}{9.2}t}$

Te-131M and Te-131 do not recirculate to blood.

Iodine as a daughter of Tellurium:

$f_1 = .95$

Source organs

1. Lungs

2. GIT

3. Thyroid: $R(t) = -0.328e^{-\frac{.693}{0.243}t} + 0.016e^{-\frac{.693}{11.3}t} + 0.312e^{-\frac{.693}{117}t}$

4. Cort bone: $R(t) = 5.7 \times 10^{-2}e^{-\frac{.693}{0.243}t} - 2.78 \times 10^{-3}e^{-\frac{.693}{11.3}t}$
 $+ 2.94 \times 10^{-3}e^{-\frac{.693}{117}t}$

5. Tra bone: $R(t) = 1.42 \times 10^{-2}e^{-\frac{.693}{0.243}t} - 6.96 \times 10^{-4}e^{-\frac{.693}{11.3}t}$
 $+ 7.35 \times 10^{-4}e^{-\frac{.693}{117}t}$

6. Kidney: $R(t) = 4.42 \times 10^{-3}e^{-\frac{.693}{0.243}t} - 2.16 \times 10^{-4}e^{-\frac{.693}{11.3}t}$
 $+ 2.28 \times 10^{-4}e^{-\frac{.693}{117}t}$

7. Muscle: $R(t) = 3.99 \times 10^{-1}e^{-\frac{.693}{0.243}t} - 1.95 \times 10^{-2}e^{-\frac{.693}{11.3}t}$
 $+ 2.06 \times 10^{-2}e^{-\frac{.693}{117}t}$

8. Liver: $R(t) = 2.56 \times 10^{-2}e^{-\frac{.693}{0.243}t} - 1.25 \times 10^{-3}e^{-\frac{.693}{11.3}t}$
 $+ 1.32 \times 10^{-3}e^{-\frac{.693}{117}t}$

$$9. \text{ Other tissue: } R(t) = 4.97 \times 10^{-1} e^{-\frac{.693}{0.243}t} - 2.43 \times 10^{-2} e^{-\frac{.693}{11.3}t} + 2.56 \times 10^{-2} e^{-\frac{.693}{117}t}$$

I-131 recirculates to blood. Absorption takes place in the UL and LL with an f_1 of .95.

Xenon as a granddaughter of Tellurium:

$$f_1 = .95$$

Source organs

1. Lungs

2. GIT

$$3. \text{ Thyroid: } R(t) = 1.0e^{-\frac{.693}{.0833}t}$$

$$4. \text{ Cort bone: } R(t) = 0.0571e^{-\frac{.693}{.0833}t}$$

$$5. \text{ Tra bone: } R(t) = 0.0143e^{-\frac{.693}{.0833}t}$$

$$6. \text{ Kidney: } R(t) = 4.43 \times 10^{-3} e^{-\frac{.693}{.0833}t}$$

$$7. \text{ Muscle: } R(t) = 0.4e^{-\frac{.693}{.0833}t}$$

$$8. \text{ Liver: } R(t) = 0.0257e^{-\frac{.693}{.0833}t}$$

$$9. \text{ Other tissue: } R(t) = 0.498e^{-\frac{.693}{.0833}t}$$

Xe-131 recirculates to blood.

Print cumulated activity ($\mu\text{Ci-days}$) at the same times as Example 1.

Again we include a discussion of how Example 2 must be coded. Follow Table 5.1 and Table B.2 to fully understand the instructions.

Information Concerning the Differential Equations

Card 1 NØ: 4 x 12 = 48 Lungs
 4 x 4 = 16 GIT
 12 compartments for Te-131M (excluding GIT and Lungs)
 12 compartments for Te-131 (excluding GIT and Lungs)
 2? compartments for I-131 (excluding GIT and Lungs)
 7 compartments for Xe-131M (excluding GIT and Lungs)

 116

TLAST, H1, EPS same as Example 1.

Card 2...Card 9 same as Example 1.

General Information for Chain of Radionuclides

Card 1 NISØ: 4
 D3: .3
 D4: .0R
 D5: .25
 NØRG: 7
 IØPT: 0
 NØRAL: not applicable, set to 0

Card 2 TAG(1): TE-131M
 TR(1): 1.25
 ICLASS(1): 0
 LAMAB(1,2): 2.
 P(1),NØRNCH(1),IØRNCH(1): not applicable
 ILAB: 0

Card 3 omitted because ILAB=0

Card 4 TAB(2): TE-131
 TR(2): 0.01736
 ICLASS(2): 0
 LAMAB(2,2): 2.
 P(2): 0.18
 NØRNCH(2): 2
 IØRNCH(2): 1
 ILAB: 0

Card 5 omitted because ILAB=0

1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48	49	50	51	52	53	54	55	56	57	58	59	60	61	62	63	64	65	66	67	68	69	70	71	72	73	74	75	76	77	78	79	80	81	82	83	84	85	86	87	88	89	90	91	92	93	94	95	96	97	98	99	100	101	102	103	104	105	106	107	108	109	110	111	112	113	114	115	116	117	118	119	120	121	122	123	124	125	126	127	128	129	130	131	132	133	134	135	136	137	138	139	140	141	142	143	144	145	146	147	148	149	150	151	152	153	154	155	156	157	158	159	160	161	162	163	164	165	166	167	168	169	170	171	172	173	174	175	176	177	178	179	180	181	182	183	184	185	186	187	188	189	190	191	192	193	194	195	196	197	198	199	200	201	202	203	204	205	206	207	208	209	210	211	212	213	214	215	216	217	218	219	220	221	222	223	224	225	226	227	228	229	230	231	232	233	234	235	236	237	238	239	240	241	242	243	244	245	246	247	248	249	250	251	252	253	254	255	256	257	258	259	260	261	262	263	264	265	266	267	268	269	270	271	272	273	274	275	276	277	278	279	280	281	282	283	284	285	286	287	288	289	290	291	292	293	294	295	296	297	298	299	300	301	302	303	304	305	306	307	308	309	310	311	312	313	314	315	316	317	318	319	320	321	322	323	324	325	326	327	328	329	330	331	332	333	334	335	336	337	338	339	340	341	342	343	344	345	346	347	348	349	350	351	352	353	354	355	356	357	358	359	360	361	362	363	364	365	366	367	368	369	370	371	372	373	374	375	376	377	378	379	380	381	382	383	384	385	386	387	388	389	390	391	392	393	394	395	396	397	398	399	400	401	402	403	404	405	406	407	408	409	410	411	412	413	414	415	416	417	418	419	420	421	422	423	424	425	426	427	428	429	430	431	432	433	434	435	436	437	438	439	440	441	442	443	444	445	446	447	448	449	450	451	452	453	454	455	456	457	458	459	460	461	462	463	464	465	466	467	468	469	470	471	472	473	474	475	476	477	478	479	480	481	482	483	484	485	486	487	488	489	490	491	492	493	494	495	496	497	498	499	500	501	502	503	504	505	506	507	508	509	510	511	512	513	514	515	516	517	518	519	520	521	522	523	524	525	526	527	528	529	530	531	532	533	534	535	536	537	538	539	540	541	542	543	544	545	546	547	548	549	550	551	552	553	554	555	556	557	558	559	560	561	562	563	564	565	566	567	568	569	570	571	572	573	574	575	576	577	578	579	580	581	582	583	584	585	586	587	588	589	590	591	592	593	594	595	596	597	598	599	600	601	602	603	604	605	606	607	608	609	610	611	612	613	614	615	616	617	618	619	620	621	622	623	624	625	626	627	628	629	630	631	632	633	634	635	636	637	638	639	640	641	642	643	644	645	646	647	648	649	650	651	652	653	654	655	656	657	658	659	660	661	662	663	664	665	666	667	668	669	670	671	672	673	674	675	676	677	678	679	680	681	682	683	684	685	686	687	688	689	690	691	692	693	694	695	696	697	698	699	700	701	702	703	704	705	706	707	708	709	710	711	712	713	714	715	716	717	718	719	720	721	722	723	724	725	726	727	728	729	730	731	732	733	734	735	736	737	738	739	740	741	742	743	744	745	746	747	748	749	750	751	752	753	754	755	756	757	758	759	760	761	762	763	764	765	766	767	768	769	770	771	772	773	774	775	776	777	778	779	780	781	782	783	784	785	786	787	788	789	790	791	792	793	794	795	796	797	798	799	800	801	802	803	804	805	806	807	808	809	810	811	812	813	814	815	816	817	818	819	820	821	822	823	824	825	826	827	828	829	830	831	832	833	834	835	836	837	838	839	840	841	842	843	844	845	846	847	848	849	850	851	852	853	854	855	856	857	858	859	860	861	862	863	864	865	866	867	868	869	870	871	872	873	874	875	876	877	878	879	880	881	882	883	884	885	886	887	888	889	890	891	892	893	894	895	896	897	898	899	900	901	902	903	904	905	906	907	908	909	910	911	912	913	914	915	916	917	918	919	920	921	922	923	924	925	926	927	928	929	930	931	932	933	934	935	936	937	938	939	940	941	942	943	944	945	946	947	948	949	950	951	952	953	954	955	956	957	958	959	960	961	962	963	964	965	966	967	968	969	970	971	972	973	974	975	976	977	978	979	980	981	982	983	984	985	986	987	988	989	990	991	992	993	994	995	996	997	998	999	1000
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Table B.2. (cont'd)

Card 6 TAG(3): I-131
 TR(3): 8.06
 ICLASS(3): D
 LAMAB(3,2): 114.
 P(3): 0.82
 NBRNCH(3): 2
 IBRNCH(3): 2
 ILAB: 1

Card 7 LAMAB(3,1) = 0
 LAMAB(3,3) = 114.
 LAMAB(3,4) = 114.

Card 8 TAG(4): XE-131M
 TR(4): 11.8
 ICLASS(4): D
 LAMAB(4,2): 114.
 P(4) = 0.0144
 NBRNCH(4): 1
 IBRNCH(4): 1
 ILAB = 0

Card 9 omitted because ILAB=0

Description of Retention

FOR TE-131M

Card 1 omitted for first nuclide in the chain

Cards 2 and 3 repeated for each organ

Cort bone

Card 2 ISORS(1): 15
 ORGNM(1): COR BONE
 ICØM(1,1): 2

Card 3 AS(1,1): 0.0183
 AS(2,1): 0.01
 TBS(1,1): 7.533×10^{-2}
 TBS(2,1): 8.663×10^{-5}

Tra bone

Card 2 ISORS(2): 16
 ORGNM(2): TRA BONE
 ICØM(1,2): 2

Card 3 AS(1,2): 4.57×10^{-3}
 AS(2,2): 0.01
 TBS(1,2): 7.533×10^{-2}
 TBS(2,2): 8.663×10^{-5}

Kidneys

Card 2 ISØRS(3): 6
ØRGNM(3): KIDNEYS
ICØM(1,3): 2

Card 3 AS(1,3): 1.42×10^{-3}
AS(2,3): 0.034
TBS(1,3): 7.533×10^{-2}
TBS(2,3): 3.013×10^{-2}

Muscle

Card 2 ISØRS(4): 12
ØRGNM(4): MUSCLE
ICØM(1,4): 2

Card 3 AS(1,4): 0.128
AS(2,4): 0.1
TBS(1,4): 7.533×10^{-2}
TBS(2,4): 3.915×10^{-2}

Liver

Card 2 ISØRS(5): 7
ØRGNM(5): LIVER
ICØM(1,5): 2

Card 3 AS(1,5): 8.23×10^{-3}
AS(2,5): 0.055
TBS(1,5): 7.533×10^{-2}
TBS(2,5): 6.794×10^{-2}

Thyroid

Card 2 ISØRS(6): 2
ØRGNM(6): THYROID
ICØM(1,6): 1

Card 3 AS(1,6): 9.14×10^{-5}
TBS(1,6): 7.533×10^{-2}

Other tissue

Card 2 ISØRS(7): 23
ØRGNM(7): ØTH TISS
ICØM(1,7): 1

Card 3 AS(1,7): 0.159
TBS(1,7): 7.533×10^{-2}

FOR TE-131

Card 1 IS(1): 0
IS(2): 0
IS(3): 0
IS(4): 0
IS(5): 0
IS(6): 0
IS(7): 0

The remainder of the cards for TE-131 are the same as for TE-131M since the retention functions are the same.

FOR I-131

Card 1 IS(1): 1
 IS(2): 1
 IS(3): 1
 IS(4): 1
 IS(5): 1
 IS(6): 1
 IS(7): 1

Cards 2 and 3 are repeated for each organ

Cort bone

Card 2 ISORS(1): 15
 ORGNM(1): COR BONE
 ICØM(3,1): 3

Card 3 AS(1,1): 5.70×10^{-2}
 AS(2,1): -2.78×10^{-3}
 AS(3,1): 2.94×10^{-3}
 TBS(1,1): 2.852
 TBS(2,1): 6.133×10^{-2}
 TBS(3,1): 5.923×10^{-3}

Tra bone

Card 2 ISORS(2): 16
 ORGNM(2): TRA BONE
 ICØM(3,2): 3

Card 3 AS(1,2): 1.42×10^{-2}
 AS(2,2): -6.96×10^{-4}
 AS(3,2): 7.35×10^{-4}
 TBS(1,2): 2.852
 TBS(2,2): 6.133×10^{-2}
 TBS(3,2): 5.923×10^{-3}

Kidneys

Card 2 ISORS(3): 6
 ORGNM(3): KIDNEYS
 ICØM(3,3): 3

Card 3 AS(1,3): 4.42×10^{-3}
 AS(2,3): -2.16×10^{-4}
 AS(3,3): 2.28×10^{-4}
 TBS(1,3): 2.852
 TBS(2,3): 6.133×10^{-2}
 TBS(3,3): 5.923×10^{-3}

Muscle

Card 2 ISORS(4): 12
 ORGNM(4): MUSCLE
 ICØM(3,4): 3

Card 3 AS(1,4): 3.99×10^{-1}
 AS(2,4): -1.95×10^{-2}
 AS(3,4): 2.05×10^{-2}
 TBS(1,4): 2.852
 TBS(2,4): 6.133×10^{-2}
 TBS(3,4): 5.923×10^{-3}

Liver

Card 2 ISØRS(5): 7
 ØRGNM(5): LIVER
 ICØM(3,5): 3

Card 3 AS(1,5): 2.56×10^{-2}
 AS(2,5): -1.25×10^{-3}
 AS(3,5): 1.32×10^{-3}
 TBS(1,5): 2.852
 TBS(2,5): 6.133×10^{-2}
 TBS(3,5): 5.923×10^{-3}

Thyroid

Card 2 ISØRS(6): 22
 ØRGNM(6): THYROID
 ICØM(3,6): 3

Card 3 AS(1,6): -0.328
 AS(2,6): 0.016
 AS(3,6): 0.312
 TBS(1,6): 2.852
 TBS(2,6): 6.133×10^{-2}
 TBS(3,6): 5.923×10^{-3}

Other tissue

Card 2 ISØRS(7): 23
 ØRGNM(7): ØTH TISS
 ICØM(3,7): 3

Card 3 AS(1,7): 4.97×10^{-1}
 AS(2,7): -2.43×10^{-2}
 AS(3,7): 2.56×10^{-2}
 TBS(1,7): 2.852
 TBS(2,7): 6.133×10^{-2}
 TBS(3,7): 5.923×10^{-3}

FOR XE-131M

Card 1 IS(1): 1
 IS(2): 1
 IS(3): 1
 IS(4): 1
 IS(5): 1
 IS(6): 1
 IS(7): 1

Card 2 and 3 are repeated for each organ

Cort bone

Card 2 ISØRS(1): 15
 ØRGNM(1): CØR BØNE
 ICØM(4,1): 1

Card 3 AS(1,1): 0.0571
 TBS(1,1): 8.319

Tra bone

Card 2 ISØRS(2): 16
 ØRGNM(2): TRA BØNE
 ICØM(4,2): 1

Card 3 AS(1,2): 0.0143
 TBS(1,2): 8.319

Kidneys

Card 2 ISORS(3): 6
 ORGNM(3): KIDNEYS
 ICDM(4,3): 1

Card 3 AS(1,3): 4.43×10^{-3}
 TBS(1,3): 8.319

Muscle

Card 2 ISORS(4): 12
 ORGNM(4): MUSCLE
 ICDM(4,4): 1

Card 3 AS(1,4): 0.4
 TBS(1,4): 8.319

Liver

Card 2 ISORS(5): 7
 ORGNM(5): LIVER
 ICDM(4,5): 1

Card 3 AS(1,5): 0.0257
 TBS(1,5): 8.319

Thyroid

Card 2 ISORS(6): 22
 ORGNM(6): THYROID
 ICDM(4,6): 1

Card 3 AS(1,6): 1.
 TBS(1,6): 8.319

Other tissue

Card 2 ISORS(7): 23
 ORGNM(7): OTH TISS
 ICDM(4,7): 1

Card 3 AS(1,7): 0.498
 TBS(1,7): 8.319

Other Tissues Transformation

Insert 12 cards all of the form

NOTOP: 1
 IOTHER(1): 15
 IOTHER(2): 16
 IOTHER(3): 6
 IOTHER(4): 12
 IOTHER(5): 7
 IOTHER(6): 22

The printout for Example 2 is given for the output to unit 10.
 Output to unit 6 is omitted because it is so lengthy. The cumulated
 activity ($\mu\text{Ci-days}$) are omitted for $T = 10 \text{ yrs.}, 20 \text{ yrs.}, 30 \text{ yrs.}, 40 \text{ yrs.},$

and 50 yrs., since these are the same as those for $T = 5$ yrs., (1826.25 days). Negative cumulated activities represent machine round-off and are in effect 0.

OUTPUT FOR EXAMPLE
TE-131H CHAIN. INTAKE BY INHALATION

PERCENT DEPOSITION							
ISOTOPE	HAIRTYPE	CLASS	LAMB	BB-RATIO	FORMCF	DEPCN	W-P
							T-D
							0.2%
TE-131H	1.2500 00	D	2.0000 00	1.0000 00	0	0	
TE-131H	1.7500 02	D	2.0000 00	1.0000 01	2	1	
TE-131H	4.0000 00	D	1.1000 02	0.2000 01	2	2	
		LAMB-S	0.0				
		LAMB-TLI	1.1000 02				
		LAMB-LTY	1.1000 02				
TE-131H	1.1000 01	D	1.1000 02	1.0000 02	1	1	

ISOTOPE	ORGAN	A'S	AND	TB'S		
TE-131H	COP BONE	1.0000-02	1.0000-02	7.5330-02	9.6630-05	
	TBA BONE	4.5700-03	1.0000-02	7.5330-02	9.6630-05	
	KIDNEYS	1.8200-03	3.0000-02	7.5330-02	3.0130-02	
	MUSCLE	1.2000-01	1.0000-01	7.5330-02	3.9150-02	
	LIVER	6.2300-03	5.5000-02	7.5330-02	6.7900-02	
	THYROID	9.1000-05	7.5330-02			
	OTH TISS	1.5500-01	7.5330-02			

ISOTOPE	ORGAN	A'S	AND	TB'S		
TE-131H	COP BONE	1.0000-02	1.0000-02	7.5330-02	9.6630-05	
	TBA BONE	4.5700-03	1.0000-02	7.5330-02	9.6630-05	
	KIDNEYS	1.8200-03	3.0000-02	7.5330-02	3.0130-02	
	MUSCLE	1.2000-01	1.0000-01	7.5330-02	3.9150-02	
	LIVER	6.2300-03	5.5000-02	7.5330-02	6.7900-02	
	THYROID	9.1000-05	7.5330-02			
	OTH TISS	1.5500-01	7.5330-02			

ISOTOPE	ORGAN	A'S	AND	TB'S			
TE-131H	COP BONE	5.7000-02	-2.7000-03	2.9000-03	2.0520 00	6.1330-02	5.9230-03
	TBA BONE	1.0200-02	-6.9600-04	7.3500-04	2.0520 00	6.1330-02	5.9230-03
	KIDNEYS	4.0200-03	-2.1600-04	2.2000-04	2.0520 00	6.1330-02	5.9230-03
	MUSCLE	1.9000-01	-1.9500-02	2.0600-02	2.0520 00	6.1330-02	5.9230-03
	LIVER	2.5600-02	-1.2500-03	1.3200-03	2.0520 00	6.1330-02	5.9230-03
	THYROID	-3.2000-01	1.6000-02	3.1400-03	2.0520 00	6.1330-02	5.9230-03
	OTH TISS	4.9700-01	-2.0300-02	2.5600-02	2.0520 00	6.1330-02	5.9230-03

ISOTOPE	ORGAN	A'S	AND	TB'S
TE-131H	COP BONE	5.7100-02	0.3190 00	
	TBA BONE	1.0330-02	0.3190 00	
	KIDNEYS	4.0390-03	0.3190 00	
	MUSCLE	4.0000-01	0.3190 00	
	LIVER	2.5700-02	0.3190 00	
	THYROID	1.0000 00	0.3190 00	
	OTH TISS	4.9800-01	0.3190 00	

TABLE 10000000

MICROORGANISMS-DAYS FROM INHALATION OF CE-1319

LUNGS

ISOTOPE	W-F	T-B	F	LYMPH
CE-1319	4.25370-33	2.99280-03	1.26160-01	1.65520-02
CE-131	2.92480-36	1.47390-34	2.19230-02	3.01370-03
I-131	4.71280-36	6.42170-36	4.32650-01	1.60750-02
CE-1319	5.74450-33	1.01930-10	2.69520-04	4.99070-07

MICROORGANISMS ENTERING
BLOOD GI

3.47040-01	1.52260-01
4.71030-02	9.64990-03
9.45950-03	1.71100-34
4.26100-06	2.17620-09

CE TRACT

ISOTOPE	STOMACH	SPL.	U.L.I.	L.L.I.
CE-1319	4.20000-33	1.71900-02	4.28920-32	4.40740-02
CE-131	4.51190-36	1.01630-03	7.80910-33	6.08420-03
I-131	2.42970-05	1.60440-35	1.28000-05	3.36730-05
CE-1319	1.70520-39	1.44740-10	1.42770-08	3.22700-08

MICROORGANISMS ENTERING
BLOOD

3.47930-02
6.06900-03
9.61110-03
1.92250-08

OTHER ORGANS

ISOTOPE	COB BONE	TFA BONE	KIDNEYS	MUSCLE	LIVER	THYROID	OTR TISS
CE-1319	1.26940-32	4.71720-03	1.42210-32	1.02290-01	2.41670-32	4.35220-05	7.04920-32
CE-131	2.12050-33	1.21750-03	2.93990-03	1.05370-02	5.16410-03	7.34270-04	1.27730-32
I-131	4.94260-04	1.71640-36	5.34310-05	4.82400-03	3.09490-04	9.99000-03	6.00590-33
CE-1319	1.83730-37	1.59940-04	1.11510-39	1.08400-06	6.46890-08	2.51710-04	1.25350-36

AFTER TRANSPORTATION

ISOTOPE	COB BONE	TFA BONE	KIDNEYS	MUSCLE	LIVER	THYROID	T. BODY
CE-1319	4.70000-03	4.69540-03	1.55940-32	4.56940-02	2.45240-32	9.35270-04	1.41510-31
CE-131	4.57200-36	4.51160-36	2.82530-03	6.28670-03	4.44470-03	1.44920-38	2.56420-32
I-131	3.10320-07	5.49240-07	1.76740-04	1.30400-04	1.41600-37	9.94730-03	1.29570-32
CE-1319	6.04770-11	4.43010-11	4.60780-10	2.44730-10	1.73150-11	2.51640-08	2.51640-36

TABLE 10000000

MICROORGANISMS-DAYS FROM INHALATION OF CE-1319

LUNGS

ISOTOPE	W-F	T-B	F	LYMPH
CE-1319	4.29170-33	2.78270-03	1.28010-01	1.84010-02
CE-131	2.92480-36	1.67820-34	2.26080-02	3.25150-03
I-131	4.71280-36	6.41000-36	7.47740-03	2.48340-02
CE-1319	5.74450-33	1.44440-10	4.17230-04	2.28330-06

MICROORGANISMS ENTERING
BLOOD GI

3.92570-01	1.52260-01
4.83100-02	9.64990-03
1.20500-02	1.71120-34
8.01990-04	2.17580-09

CE TRACT

ISOTOPE	STOMACH	SPL.	U.L.I.	L.L.I.
CE-1319	4.20000-33	1.71970-02	4.38790-32	5.14320-02
CE-131	4.51200-36	1.01640-03	7.81630-33	6.02140-03
I-131	2.42970-05	1.60450-35	1.28500-05	3.42130-05
CE-1319	1.70540-39	1.44740-10	1.58500-08	5.87640-08

MICROORGANISMS ENTERING
BLOOD

3.47930-02
6.06990-03
1.03100-02
1.92280-08

OTHER ORGANS

ISOTOPE	COB BONE	TFA BONE	KIDNEYS	MUSCLE	LIVER	THYROID	OTR TISS
CE-1319	1.97830-02	1.05480-02	2.52940-02	1.54140-01	4.26600-02	6.10760-05	1.06250-01
CE-131	3.50640-33	1.92040-03	4.59440-03	2.83680-02	7.74970-03	1.10950-05	1.93040-32
I-131	1.15420-33	2.87430-04	4.94170-05	4.07570-03	5.17900-04	6.85010-02	1.00430-32
CE-1319	5.58850-07	1.39440-07	4.33500-08	3.91490-06	2.51530-07	9.78730-04	4.87410-36

AFTER TRANSPORTATION

ISOTOPE	COB BONE	TFA BONE	KIDNEYS	MUSCLE	LIVER	THYROID	T. BODY
CE-1319	7.55530-03	7.42140-03	2.43510-02	7.08230-02	1.71740-02	1.36450-07	2.13290-01
CE-131	1.37300-03	1.16490-03	4.42400-03	1.28680-02	4.75140-03	2.87820-08	3.07440-02
I-131	2.04910-06	5.41460-07	1.29200-07	1.09770-05	4.72240-07	6.85750-02	2.01620-02
CE-1319	2.54880-10	1.40010-10	2.64230-11	1.17460-09	4.75130-11	9.78850-06	9.78850-36

24-11-1963 (1)

MICROCUBIES-DAYS FROM IRRADIATION OF TE-131a

LUNGS					MICROCUBIES ENTERING BLOOD
ISOTOPE	S-P	T-B	P	LIVER	GI
TE-131a	6.29370-03	2.09270-03	1.20010-01	1.00020-02	3.92570-01
TE-131	2.92000-00	1.47020-00	2.20000-02	1.35150-03	4.03100-02
I-131	6.71290-06	6.81000-06	7.47000-03	2.00550-03	1.20500-02
IE-131a	5.70650-11	1.00000-10	6.37770-06	2.29490-06	4.00200-06

II TRACT

ISOTOPE	STOMACH	S.I.	U.L.I.	L.L.I.	MICROCUBIES ENTERING BLOOD
TE-131a	6.20000-03	1.71970-02	6.30790-02	5.10360-02	3.67930-02
TE-131	6.51200-00	3.03000-03	7.41630-03	6.82050-03	6.06990-03
I-131	2.42000-05	1.00050-05	3.32500-05	3.92160-05	1.03190-02
IE-131a	1.70000-04	1.00100-03	1.50500-00	5.00700-00	3.92200-00

OTHER ORGANS

ISOTOPE	COB BONE	TBA BONE	KIDNEYS	MUSCLE	LIVER	THYROID	OTH TISS
TE-131a	2.01230-02	1.00070-02	2.50170-02	1.50000-01	6.33000-02	6.20170-05	1.07090-01
TE-131	1.65620-03	1.06190-03	6.69150-03	2.00590-02	7.47000-03	1.12660-05	1.95990-02
I-131	1.40130-03	6.00050-00	1.27290-06	1.15000-02	7.37300-04	1.09010-01	1.02000-02
IE-131a	1.13400-04	2.00000-07	1.20100-00	4.30000-04	4.30000-07	2.00000-05	1.20000-05

AFTER TRANSFORMATION

ISOTOPE	COB BONE	TBA BONE	KIDNEYS	MUSCLE	LIVER	THYROID	T. BODY
TE-131a	7.70710-03	7.71270-03	2.00500-02	7.22090-02	3.77770-02	1.30500-07	2.16570-01
TE-131	1.00000-03	1.00100-03	6.41730-03	1.11100-02	6.00310-03	2.51600-06	1.01000-02
I-131	6.12330-06	1.97770-07	2.00150-07	2.67330-05	2.02000-07	1.00000-01	2.00000-02
IE-131a	5.50000-10	1.00000-10	6.00000-11	1.00000-09	1.00000-10	2.00000-05	2.00000-05

24-11-1963 (2)

MICROCUBIES-DAYS FROM IRRADIATION OF TE-131a

LUNGS					MICROCUBIES ENTERING BLOOD
ISOTOPE	S-P	T-B	P	LIVER	GI
TE-131a	6.29370-03	2.09270-03	1.20010-01	1.00020-02	3.92570-01
TE-131	2.92000-00	1.47020-00	2.20000-02	1.35150-03	4.03100-02
I-131	6.71290-06	6.81000-06	7.47000-03	2.00550-03	1.20500-02
IE-131a	5.70650-11	1.00000-10	6.37770-06	2.29490-06	4.00200-06

II TRACT

ISOTOPE	STOMACH	S.I.	U.L.I.	L.L.I.	MICROCUBIES ENTERING BLOOD
TE-131a	6.20000-03	1.71970-02	6.30790-02	5.10360-02	3.67930-02
TE-131	6.51200-00	3.03000-03	7.41630-03	6.82050-03	6.06990-03
I-131	2.42000-05	1.00050-05	3.32500-05	3.92160-05	1.03190-02
IE-131a	1.70000-04	1.00100-03	1.50500-00	5.00700-00	3.92200-00

OTHER ORGANS

ISOTOPE	COB BONE	TBA BONE	KIDNEYS	MUSCLE	LIVER	THYROID	OTH TISS
TE-131a	2.01230-02	1.00070-02	2.50170-02	1.50000-01	6.33000-02	6.20170-05	1.07090-01
TE-131	1.65620-03	1.06190-03	6.69150-03	2.00590-02	7.47000-03	1.12660-05	1.95990-02
I-131	1.40130-03	6.00050-00	1.27290-06	1.15000-02	7.37300-04	1.09010-01	1.02000-02
IE-131a	1.13400-04	2.00000-07	1.20100-00	4.30000-04	4.30000-07	2.00000-05	1.20000-05

AFTER TRANSFORMATION

ISOTOPE	COB BONE	TBA BONE	KIDNEYS	MUSCLE	LIVER	THYROID	T. BODY
TE-131a	7.70710-03	7.71270-03	2.00500-02	7.22090-02	3.77770-02	1.30500-07	2.16570-01
TE-131	1.00000-03	1.00100-03	6.41730-03	1.11100-02	6.00310-03	2.51600-06	1.01000-02
I-131	6.12330-06	1.97770-07	2.00150-07	2.67330-05	2.02000-07	1.00000-01	2.00000-02
IE-131a	5.50000-10	1.00000-10	6.00000-11	1.00000-09	1.00000-10	2.00000-05	2.00000-05

T-1-14-1311

MICROBODIES-DAYS FROM INHALATION OF T-1-1311

LUNGS

ISOTOPE	B-F	T-B	F	LYMPH
T-1311	4.24370-33	2.76270-03	1.40000-01	1.96020-02
T-131	2.62400-06	1.67620-06	2.26000-02	3.75150-03
I-131	4.71290-06	4.31000-06	7.47000-03	2.00450-03
IS-1311	5.76450-11	1.06690-10	4.17770-06	2.29490-06

MICROBODIES ENTERING
BLOOD
GI

3.92470-01	1.52260-01
4.83100-02	9.94990-03
1.20540-02	1.71120-06
0.00200-06	2.17500-09

GI TRACT

ISOTOPE	SCYRACH	S.I.	D.L.I.	L.L.I.
T-1311	4.22000-33	1.73970-02	4.34790-02	4.16300-02
T-131	4.51200-06	1.03690-03	7.91630-03	9.02450-03
I-131	2.92900-05	1.80650-05	1.32540-05	1.92100-05
IS-1311	1.79540-09	1.66100-10	1.50500-05	5.09700-08

MICROBODIES ENTERING
BLOOD

3.67930-02
4.06990-03
1.03190-02
3.42200-08

OTHER ORGANS

ISOTOPE	COR BONE	TIB BONE	KIDNEYS	MUSCLE	LIVER	THYROID	OTH TISS
T-1311	2.03230-32	1.08670-02	2.56170-02	1.50640-01	0.33460-02	6.20170-05	1.07090-01
T-131	1.65420-03	1.96990-03	4.69150-03	2.00490-02	7.87400-03	1.12660-05	1.65990-02
I-131	1.75510-03	4.37890-06	1.35950-06	1.22030-02	7.87400-06	1.03050-01	1.52610-02
IS-1311	1.22710-06	1.22350-07	9.98660-08	9.01670-08	5.79320-07	2.25420-05	1.12260-05

AFTER TRANSFORMATION

ISOTOPE	COR BONE	TIB BONE	KIDNEYS	MUSCLE	LIVER	THYROID	T. BODY
T-1311	7.76710-03	7.71270-03	2.40500-02	7.22090-02	3.77770-02	1.10540-07	2.16570-01
T-131	1.03000-03	1.40100-03	4.51730-03	1.37210-02	4.06310-03	2.51620-00	1.93440-02
I-131	4.51670-05	1.56190-07	2.42390-07	2.06910-05	2.02200-07	1.03040-01	3.06350-02
IS-1311	4.76410-10	4.16610-10	4.08620-11	2.50660-09	1.55660-10	2.25350-05	2.25350-05

T-1-14-1311

MICROBODIES-DAYS FROM INHALATION OF T-1-1311

LUNGS

ISOTOPE	B-F	T-B	F	LYMPH
T-1311	4.24370-33	2.76270-03	1.40000-01	1.96020-02
T-131	2.62400-06	1.67620-06	2.26000-02	3.75150-03
I-131	4.71290-06	4.31000-06	7.47000-03	2.00450-03
IS-1311	5.76450-11	1.06690-10	4.17770-06	2.29490-06

MICROBODIES ENTERING
BLOOD
GI

3.92470-01	1.52260-01
4.83100-02	9.94990-03
1.20540-02	1.71120-06
0.00200-06	2.17500-09

GI TRACT

ISOTOPE	SCYRACH	S.I.	D.L.I.	L.L.I.
T-1311	4.22000-33	1.73970-02	4.34790-02	4.16300-02
T-131	4.51200-06	1.03690-03	7.91630-03	9.02450-03
I-131	2.92900-05	1.80650-05	1.32540-05	1.92100-05
IS-1311	1.79540-09	1.66100-10	1.50500-05	5.09700-08

MICROBODIES ENTERING
BLOOD

3.67930-02
4.06990-03
1.03190-02
3.42200-08

OTHER ORGANS

ISOTOPE	COR BONE	TIB BONE	KIDNEYS	MUSCLE	LIVER	THYROID	OTH TISS
T-1311	2.03230-32	1.08670-02	2.56170-02	1.50640-01	0.33460-02	6.20170-05	1.07090-01
T-131	1.65420-03	1.96990-03	4.69150-03	2.00490-02	7.87400-03	1.12660-05	1.65990-02
I-131	1.75510-03	4.37890-06	1.35950-06	1.22030-02	7.87400-06	1.03050-01	1.52610-02
IS-1311	1.22710-06	1.22350-07	9.98660-08	9.01670-08	5.79320-07	2.25420-05	1.12260-05

AFTER TRANSFORMATION

ISOTOPE	COR BONE	TIB BONE	KIDNEYS	MUSCLE	LIVER	THYROID	T. BODY
T-1311	7.76710-03	7.71270-03	2.40500-02	7.22090-02	3.77770-02	1.10540-07	2.16570-01
T-131	1.03000-03	1.40100-03	4.51730-03	1.37210-02	4.06310-03	2.51620-00	1.93440-02
I-131	4.51670-05	1.56190-07	2.42390-07	2.06910-05	2.02200-07	1.03040-01	3.06350-02
IS-1311	4.76410-10	4.16610-10	4.08620-11	2.50660-09	1.55660-10	2.25350-05	2.25350-05

74 1.0220 73

MICROBILES-DAYS FROM IRRADIATION OF TE-1319

LEADS

DATE	S-P	T-B	F	LIVED
TE-1319	8.25970-13	2.74275-03	1.20010-01	1.00020-02
TE-131	2.42000-04	1.47020-04	2.20300-02	1.35130-03
T-131	8.71290-04	8.43000-06	7.47000-03	2.40550-03
TE-1319	5.70000-11	1.00000-10	0.37770-00	2.29000-00

MICROBILES ENTERING BLOOD GI

3.42570-01	1.52200-01
0.83100-02	0.80000-03
1.20500-02	1.71120-04
0.00200-00	2.17500-09

DE TRACT

DATE	STOMACH	S-I-	S-L-I-	S-L-L-I-
TE-1319	6.23000-03	1.73070-02	4.30700-02	5.16300-02
TE-131	2.71200-00	1.03000-03	7.01000-03	0.02000-03
T-131	1.02000-05	1.03000-05	1.32500-05	1.32100-05
TE-1319	1.00000-00	1.00000-10	1.50000-00	5.00700-00

MICROBILES ENTERING SLO-70

1.47000-02
6.00000-03
1.21100-02
1.02200-00

CPDS-190855

DATE	CC9 9000	T9A 9000	K10000	90000	LIVED	T90000	JTR 1000
TE-1319	2.31200-02	1.00000-02	2.50170-02	1.50000-01	0.03000-02	0.20170-04	1.07000-01
TE-131	1.40000-03	1.00000-03	0.00100-03	2.00000-02	7.07000-03	1.12000-00	1.05000-02
T-131	1.70000-03	0.37000-04	1.35000-04	1.22000-02	7.67000-04	1.03000-01	1.52000-02
TE-1319	1.22000-04	1.22000-07	1.00000-00	0.01000-00	0.70000-07	2.25000-00	1.12200-05

LETER TRANSFORMATION

DATE	CC9 9000	T9A 9000	K10000	90000	LIVED	T90000	JTR 1000
TE-1319	7.70000-03	7.71270-03	2.00000-02	7.22000-02	1.77770-02	1.10500-03	2.10570-01
TE-131	1.40000-03	1.00000-03	0.51770-03	1.31000-02	0.90000-03	2.51000-04	1.00000-02
T-131	0.50000-05	-1.50000-07	2.07000-07	2.00000-05	-2.01000-07	1.00000-01	1.00000-02
TE-1319	-5.00000-10	4.70000-10	0.30000-11	2.50700-09	-1.50000-10	2.25000-05	2.25000-05

Example 3

Radionuclide - I-132

Half life - 2.38 hours

Mode of intake - ingestion

$$f_1 = 0.95$$

Source organs

1. GIT

$$2. \text{ Thyroid: } R(t) = -0.328e^{-\frac{.693}{0.24265}t} + 0.016e^{-\frac{.693}{11.3235}t} \\ + 0.312e^{-\frac{.693}{117.378}t}$$

$$3. \text{ Total body: } R(t) = 0.9973e^{-\frac{.693}{0.24265}t} - 0.04873e^{-\frac{.693}{11.3235}t} \\ + 0.05143e^{-\frac{.693}{117.378}t}$$

Print cumulated activity ($\mu\text{Ci-days}$) for same times as Example 1.

A description of the input cards is given below. Follow Table 5.1 and Table B.3 to fully understand the instructions.

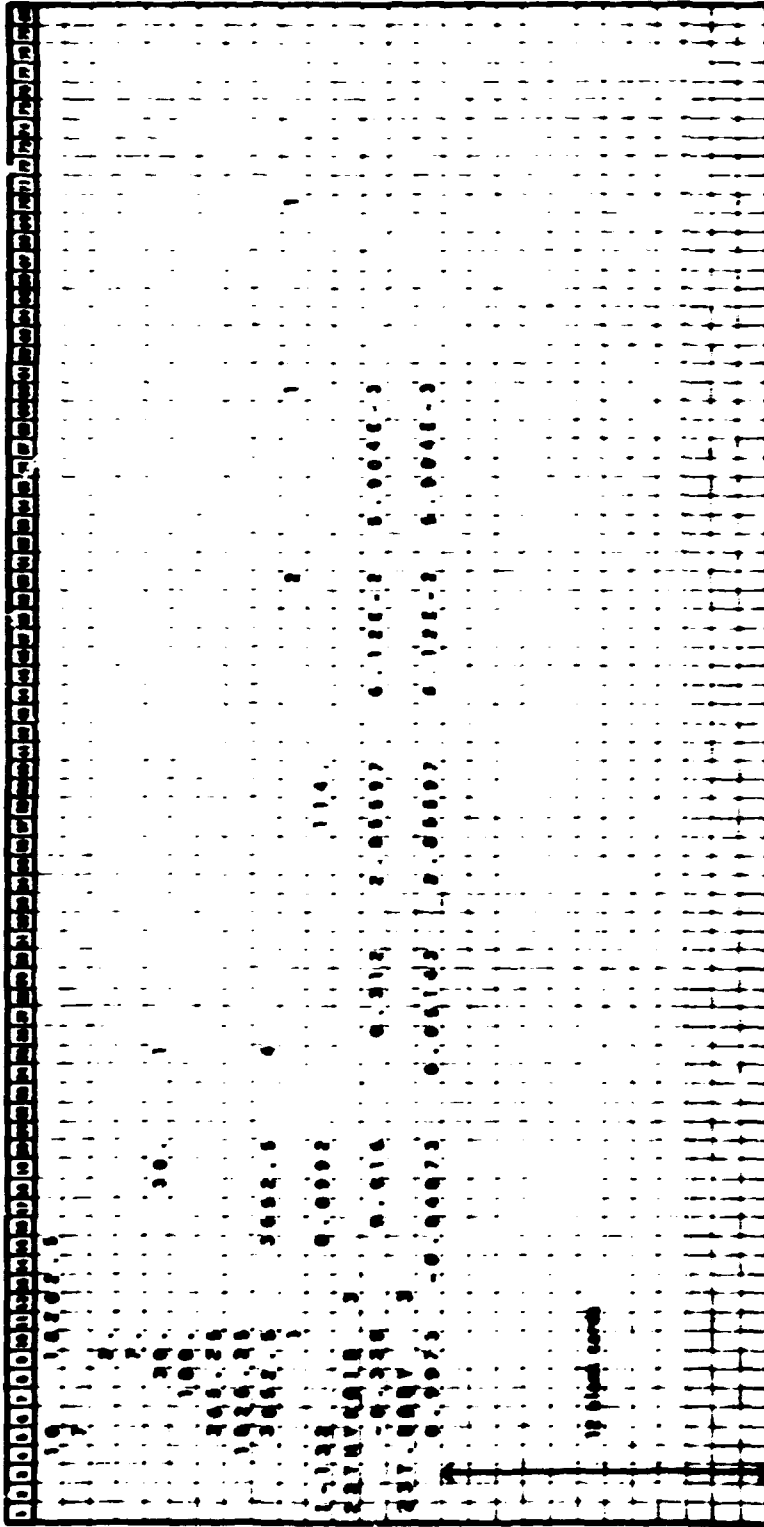
Information Concerning the Differential Equations

<u>Card 1</u>	NO:	4	GIT
		6	Thyroid and Total body
		10	

TLAST, H1, EPS same as Example 1.

Card 2...Card 9 same as Example 1.

Table 0.2. 1-120; Inside by Impaction



General Information for Chain of Radioisotopes

Card 1 NISD: 4
 D3: }
 D4: } not applicable
 D5: }
 NORG: 2
 IPPT: 1
 NORAL: 1

Card 2 TAG(1): I-132
 TR(1): 0.0992
 ICLASS(1): not applicable
 LAMAB(1,2): 114.
 P(1), NBRNCH(1), IBFNCH(1): not applicable
 ILAB: 0

Card 3 omitted because ILAB=0.

Description of Retention

Card 1 omitted for first nuclide in the chain.

Card 2 and 3 repeated for each organ.

Thyroid

Card 2 ISORS(1): 22
 ORGNM(1): THYROID
 ICOM(?,1): 3

Card 3 AS(1,1): -0.328
 AS(2,1): 0.016
 AS(3,1): 0.312
 TBS(1,1): 2.85597
 TBS(2,1): 6.12×10^{-2}
 TBS(3,1): 5.904×10^{-3}

Total body

Card 2 ISORS(2): 23
 ORGNM(2): T. BODY
 ICOM(1,2): 3

Card 3 AS(1,2): 0.9973
 AS(2,2): -0.04873
 AS(3,2): 0.05143
 TBS(1,2): 2.85597
 TBS(2,2): 6.12×10^{-2}
 TBS(3,2): 5.904×10^{-3}

Other Tissues Transformation

Insert 12 blank cards.

Printout to both unit 6 and unit 10, is included for Example 3. Printout for cumulated activity ($\mu\text{Ci-days}$) for T greater than 7 days has been omitted since there is no change in the cumulated activity after 7 days.

EXAMPLE 3
STATE BY INGESTION

DL POINT INTERVALS DEFINED = 7 DL READ = 7
POINTS SELECTED IN POINT INTERVAL = 100000 STATE BY INGESTION
T
2.0000000000 00 0.0 0
7.0000000000 01 0.0 0
1.0000000000 02 1.0000000000 00 1
1.0000000000 02 0.0 0
1.5000000000 02 0.0 0
1.0000000000 03 0.0 0
1.5000000000 03 1.5000000000 00 0

DL 10
DL 21
DL 0
DL 50 1.0000000000 00
DL 100 1.0000000000 00
DL 1.0000000000

ENTRIES OF A	NUM	COL
-0.2400000 C2	1	1
0.2400000 C2	2	1
-0.1200000 C1	2	2
0.0000000 C1	3	2
-0.1800000 C1	3	3
0.1800000 C1	4	3
-0.1000000 C1	4	4
0.0	5	1
-0.3733200 C2	5	2
0.0	5	3
0.0	5	4
-0.2455970 C1	5	5
0.0	6	1
0.1024000 C1	6	2
0.0	6	3
0.0	6	4
-0.6120000 C1	6	5
0.0	7	1
0.3550000 C2	7	2
0.0	7	3
0.0	7	4
-0.5910000 C2	7	7
0.0	8	1
0.1110000 C1	8	2
0.0	8	3
0.0	8	4
-0.2455970 C1	8	5
0.0	9	1
-0.5555220 C1	9	2
0.0	9	3
0.0	9	4
-0.1200000 C1	9	7
0.0	10	1
-0.5400000 C1	10	2
0.0	10	3
0.0	10	4
-0.5400000 C2	10	10

REPLAC = -1 FROM SCIPZ AT T = 0.0
 ERROR TEST FAILED WITH ABS(W) = 0.001

W HAS BEEN REDUCED TO 0.000000000000000000 AND STEP WILL BE RETRIED

T= 0.000000000000000000	Y FOLLOWS	4.09910119710-03	4.18269736800-03	9.5718675070-04	-2.31687261510-02	1.57830070690-03
3.227121162 90-02	6.09910119710-03	4.18269736800-03	9.5718675070-04	-2.31687261510-02	1.57830070690-03	
3.102021722 30-02	7.08856820850-02	-4.80691209050-03	5.11336865320-03			
T= 0.000000000000000000	Y FOLLOWS	4.09910119710-03	4.18269736800-03	9.57523770810-04	-2.31687250660-02	1.57830070690-03
3.227121162 90-02	6.09910119710-03	4.18269736800-03	9.57523770810-04	-2.31687250660-02	1.57830070690-03	
3.102021722 30-02	7.08856820850-02	-4.80691209050-03	5.11337896960-03			
T= 0.000000000000000000	Y FOLLOWS	4.09910119710-03	4.18269736800-03	9.57518673890-04	-2.31687261510-02	1.57830070690-03
3.227121162 90-02	6.09910119710-03	4.18269736800-03	9.57518673890-04	-2.31687261510-02	1.57830070690-03	
3.102021722 30-02	7.08856820850-02	-4.80691209050-03	5.11336465000-03			
T= 0.000000000000000000	Y FOLLOWS	4.09910119710-03	4.18269736800-03	9.57518675070-04	-2.31687261510-02	1.57830070690-03
3.227121162 90-02	6.09910119710-03	4.18269736800-03	9.57518675070-04	-2.31687261510-02	1.57830070690-03	
3.102021722 30-02	7.08856820850-02	-4.80691209050-03	5.11334865320-03			
T= 0.000000000000000000	Y FOLLOWS	4.09910119710-03	4.18269736800-03	9.57518675070-04	-2.31687261510-02	1.57830070690-03
3.227121162 90-02	6.09910119710-03	4.18269736800-03	9.57518675070-04	-2.31687261510-02	1.57830070690-03	
3.102021722 30-02	7.08856820850-02	-4.80691209050-03	5.11334865320-03			
T= 0.000000000000000000	Y FOLLOWS	4.09910119710-03	4.18269736800-03	9.57518675070-04	-2.31687261510-02	1.57830070690-03
3.227121162 90-02	6.09910119710-03	4.18269736800-03	9.57518675070-04	-2.31687261510-02	1.57830070690-03	
3.102021722 30-02	7.08856820850-02	-4.80691209050-03	5.11336865320-03			
T= 0.000000000000000000	Y FOLLOWS	4.09910119710-03	4.18269736800-03	9.57518675070-04	-2.31687261510-02	1.57830070690-03
3.227121162 90-02	6.09910119710-03	4.18269736800-03	9.57518675070-04	-2.31687261510-02	1.57830070690-03	
3.102021722 30-02	7.08856820850-02	-4.80691209050-03	5.11336865320-03			
T= 0.000000000000000000	Y FOLLOWS	4.09910119710-03	4.18269736800-03	9.57518675070-04	-2.31687261510-02	1.57830070690-03
3.227121162 90-02	6.09910119710-03	4.18269736800-03	9.57518675070-04	-2.31687261510-02	1.57830070690-03	
3.102021722 30-02	7.08856820850-02	-4.80691209050-03	5.11336865320-03			
T= 0.000000000000000000	Y FOLLOWS	4.09910119710-03	4.18269736800-03	9.57518675070-04	-2.31687261510-02	1.57830070690-03
3.227121162 90-02	6.09910119710-03	4.18269736800-03	9.57518675070-04	-2.31687261510-02	1.57830070690-03	
3.102021722 30-02	7.08856820850-02	-4.80691209050-03	5.11334865320-03			

REPLAC

ISOTOPE	HAIFLIFE	CLASS	LABOR	DR. RATIO	EFFICIENCY	ISORCM
I-132	9.5200-02	0	1.1400 02	1.0000 00	0	0

ISOTOPE	ORGAN	A'S AND TB'S					
I-132	THYROID	-3.2400-01	1.6000-02	3.1200-01	2.0560 00	6.1200-02	5.9000-03
	I.BODY	9.9730-01	-0.0730-02	5.1430-02	2.0560 00	6.1200-02	5.9000-03

T = 2.0000 00

MICROCURIE-DAYS FROM INGESTION OF I-132

GI TRACT					MICROCURIES ENTERING BLOOD
ISOTOPE	STOMACH	S.I.	U.L.I.	L.L.I.	
I-132	3.2270-02	4.0990-03	8.14270-03	9.575 00-04	6.95300-01

OTHER ORGANS

ISOTOPE	THYROID	I.BODY
I-132	9.02970-03	7.07520-02

T = 7.0000 00

MICROCURIE-DAYS FROM INGESTION OF I-132

GI TRACT					MICROCURIES ENTERING BLOOD
ISOTOPE	STOMACH	S.I.	U.L.I.	L.L.I.	
I-132	3.2270-02	4.0990-03	8.14270-03	9.575 00-04	6.95300-01

OTHER ORGANS

ISOTOPE	THYROID	I.BODY
I-132	9.02990-03	7.07520-02

APPENDIX C

JOB CONTROL LANGUAGE STATEMENTS

The following Job Control Language (JCL) cards have been used to run TIME on the IBM System/360 machines. After debugging we have avoided the costly compilation steps by obtaining hex decks and bypassing the compilation. If the user wishes to obtain the printout on unit 6, omit the DD card associated with it. If output to unit 7, 14, or 10 is to be suppressed, replace the DD parameters with DUMMY as was done in the case of unit 6.

```

1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30 31 32 33 34 35 36 37 38 39 40 41 42 43 44 45 46 47 48 49 50 51 52 53 54 55 56 57 58 59 60 61 62 63 64 65 66 67 68 69 70 71 72 73 74 75 76 77 78 79 80
// JOB CARD
// EXEC ASMFCL.PARM.ASM='LOAD'
// ASM.SYSIN DD *
//
//   A.ZED.SOURCE.DECK
//
// *
// EXEC ASMFCL.PARM.ASM='LOAD'
// ASM.SYSIN DD *
//
//   V.PAY.SOURCE.DECK
//
// *
// EXEC FORTNCL0.PARM.FORT='IRRE'
// PARM.CB='SB=51,SI=50,EU=-1,DUMP=1'
// RESIDM.SB=320K
// FORT.SYSIN DD *
//
//   FORT.RAN.SOURCE.DECK
//
// *
// DD.FT10.F001 DD DUMMY          SUPPRESS OUTPUT ON UNIT 6
// DD.FT07.F001 DD SYSOUT=B       PUNCHED OUTPUT
// DD.FT14.F001 DD UNIT=SYSDA,DSN=ABUC1DAY,DISP=(NEW,PASS)
// DCB=(LRECFM=VBIS,LRECL=180,BLKSIZE=1804)

```

