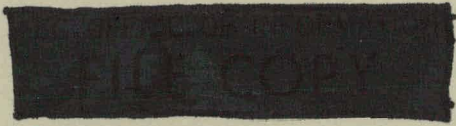


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DACRIN - A COMPUTER PROGRAM
 FOR CALCULATING ORGAN DOSE
 FROM ACUTE OR CHRONIC RADIONUCLIDE
 INHALATION

MASTER

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Pacific Northwest Laboratories
Richland, Washington 99352

DECEMBER 1974

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December 1974

OCCUPATIONAL AND ENVIRONMENTAL SAFETY DEPARTMENT

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TABLE OF CONTENTS

	<u>Page</u>
INTRODUCTION	1
SUMMARY	2
Mathematical Models and Methods	3
REFERENCES	7
APPENDIX A - DESCRIPTION OF MATHEMATICAL MODELS	A-1
APPENDIX B - COMPUTER CODE	B-1
APPENDIX C - CODE FLOW DIAGRAM	C-1
APPENDIX D - DATA LIBRARIES	D-1
APPENDIX E - PROGRAM LISTING	E-1
APPENDIX F - INPUT PREPARATION	F-1
APPENDIX G - SAMPLE PROBLEMS	G-1

DACRIN - A COMPUTER PROGRAM FOR CALCULATING
ORGAN DOSE FROM ACUTE OR CHRONIC RADIONUCLIDE INHALATION

J. R. Houston, D. L. Streng, and E. C. Watson

INTRODUCTION

The lung model, proposed by the ICRP Task Group on Lung Dynamics,^(1,2) permits a more realistic calculation of radiation dose to the human respiratory tract from the inhalation of radioactive aerosols than does the initial ICRP lung model.⁽³⁾ As is the case with most models the more closely the model approaches reality the more complex the mathematics describing the model become. The present case is no exception. This quantum increase in complexity has removed the calculation of the dose to an organ following inhalation of a radionuclide from the convenient realm of the slide rule and simple "desk top" calculator. This is particularly true when calculating the radiation dose to internal organs other than the lung. The computer program presented here has been designed to quickly and easily compute the effective dose to the respiratory tract and other organs and tissues of interest following either acute or chronic inhalation of a radioactive aerosol. A detailed description of the mathematical models used in the code is presented in Appendix A.

SUMMARY

The computer program, DACRIN, permits rapid and consistent estimates of the effective radiation dose to the human respiratory tract and other organs resulting from the inhalation of radioactive aerosols. The program is an outgrowth of the development of a mathematical model for the organ dose following the basic precepts of the ICRP Task Group on Lung Dynamics and a simple exponential model for retention by an organ of interest. As an adjunct to the organ dose model, mathematical models describing atmospheric dispersion have been included for the purpose of evaluating doses resulting from either accidental or chronic atmospheric releases of radionuclides.

The program will calculate the effective radiation dose to any of 18 organs and tissues from inhalation of any one or combination of radionuclides considered by the ICRP. A maximum of 10 organs may be selected for any one case (run). In addition, up to five multiple intake intervals and 10 time intervals measured from the last intake, may be selected for each case. Organ doses from inhalation can be calculated by specifying either the quantity of a radionuclide inhaled or the quantity released to the atmosphere. In the latter case, the duration of release, the release height, wind speed, atmospheric dispersion parameters and downwind distance at which the dose is to be calculated must also be specified. As many as 10 distances may be specified for each case.

Input to the code, in its simplest form, consists of a few program control variables, the duration of inhalation exposure, ventilation rate, the time interval within which the dose is delivered, the organs of interest, the quantity of the radionuclide inhaled, its solubility class and its particle size. Input to the code in its most complex form, results from invoking an atmospheric dispersion model. It is then necessary to input a number of additional parameters, the exact number being determined by the particular atmospheric dispersion model selected for analysis. Data libraries accessed by the code provide most of the basic data required to complete the calculations.

Output of the code consists of the effective radiation dose to the selected organs at selected time intervals, for each radionuclide inhaled as indicated by the input.

Models and Methods

The model of the respiratory tract adopted by the Task Group on Lung Dynamics forms the general basis for the mathematical models developed to calculate the dose from the inhalation of radionuclides.⁽¹⁾ In this model, the respiratory tract is divided into three regions, the nasopharyngeal (NP), the tracheobronchial (TB), and the pulmonary (P). The schematic representation of the respiratory tract used in the development of the mathematical model for the deposition and clearance of inhaled radionuclides is shown in Figure 1. Deposition is assumed to vary with the aerodynamic properties of the aerosol distribution and is described by the three parameters D_3 , D_4 , and D_5 . These parameters represent the fraction of the inhaled material, Q_I , initially deposited in the NP, TB, and P regions, respectively. Each of the three regions of deposition are further subdivided into two or more subcompartments. Each subcompartment represents the fraction of material initially in a compartment that is subject to a particular clearance process. This fraction is represented by f_k , where k indicates the clearance pathway. The quantity of material in the TB region, for example, cleared by process (c) is then represented by the product $f_c D_4 Q_I$. Values of the (f_k) and the clearance half times T_k for each clearance process for the three solubility classes of aerosols used in the code are those suggested by the ICRP,⁽²⁾ see Appendix A, Table A-5. Values of the deposition fractions D_3 , D_4 , and D_5 as function of activity median aerodynamic diameter in the form of a graph have been published.⁽¹⁾ Routines to generate these values directly from the AMAD have been included in the code and yield essentially the same values as those presented by the Task Group for the range of particle size distributions considered by that group.

The respiratory tract model was incorporated by Voillequè into a simple metabolic model for acute inhalation exposures.⁽⁴⁾ The model was programmed into a computer code called AERIN. In this model, transport of a radionuclide from the respiratory tract lymphatic system and GI tract

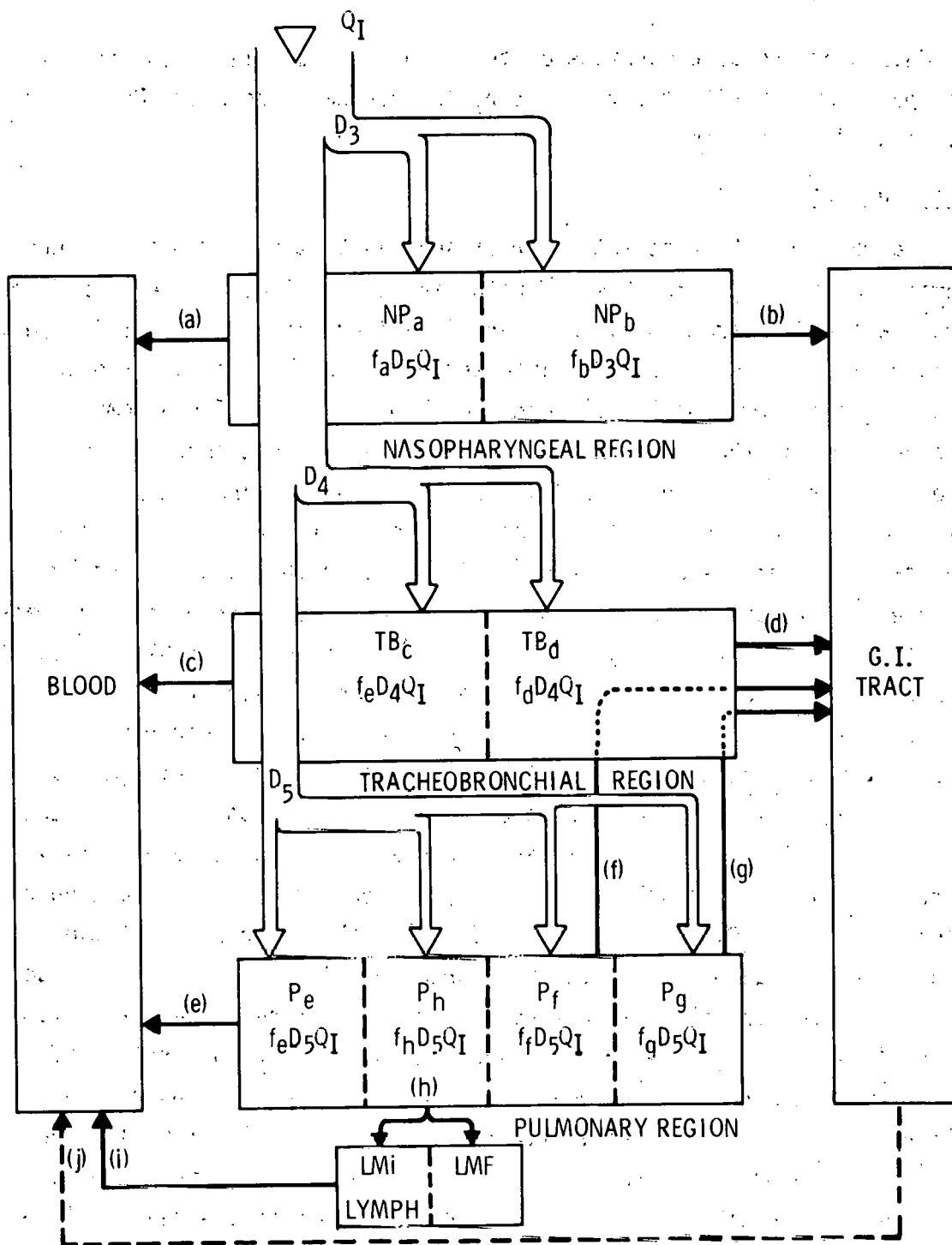


FIGURE 1. Schematic Diagram of the Task Group Lung Model

to other organs and tissues where significant accumulations of the inhaled radionuclides occur, is assumed to take place via the blood. This translocation from the respiratory tract and lymphatic system to the blood has been described in some detail by the Task Group. Of the material clearing from the respiratory tract through the GI tract, a constant fraction, f_1 is assumed to be taken up by the blood. That moving to the n^{th} organ or tissue is assumed to be a constant fraction, f_{2n} of the amount entering the blood stream at any time t . Once in the n^{th} organ, the activity is assumed to clear the organ (and the body) at a constant rate, λ_n . Voillequè's program, AERIN⁽⁴⁾, calculates the quantity of a radionuclide present in and the dose received by organs of interest as a function of time following acute exposures.

The present code extends previous codes based on the Task Group Lung Model to include calculating organ doses resulting from chronic inhalation exposure. A schematic presentation of the DACRIN metabolic model is shown in Figure 2. A model for the dose to the GI tract from radionuclides moving through it is not included in the present version of the code, although some provisions have been made in the code for the eventual addition of a GI tract dose model.

The contribution to the pulmonary lung dose from daughter nuclides formed there is computed indirectly by utilizing weighted values of the effective energy emitted by the daughter nuclides in the chain. Weighted values were calculated for each of the decay chains tabulated by the ICRP^(3,5) for residence half times of 1 day, 50 days and 500 days. These values were included in the organ data library.

Normalized air concentrations at desired distances as calculated by any atmospheric dispersion model can be used as input to DACRIN. The bivariate normal distribution model, however, was incorporated in the code because of its wide use and acceptance. As part of the input to the bivariate model, the standard deviation of the cloud concentration in the crosswind lateral and vertical directions is needed. Of the several methods in common use for estimating these standard deviations, three have been selected and programmed, i.e., Sutton's parameters, the Hanford equations, and values

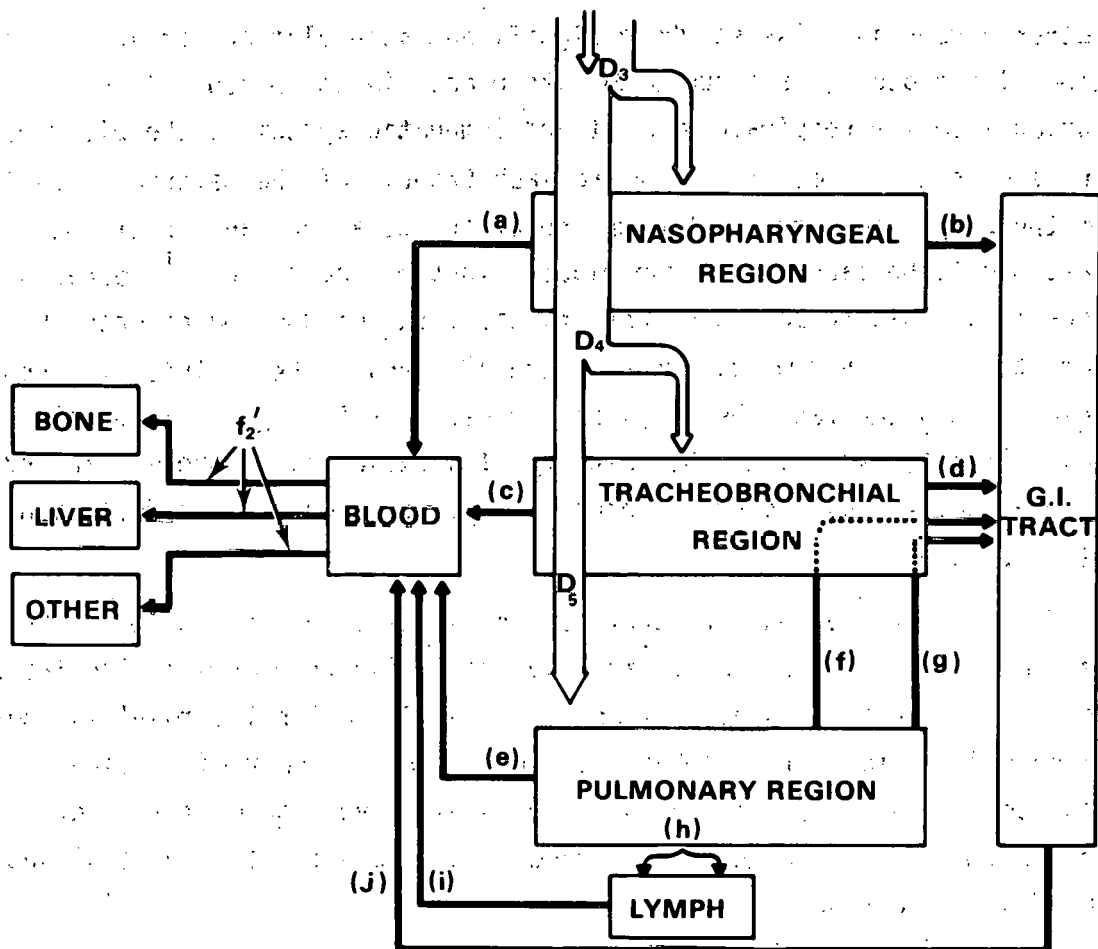


FIGURE 2: Schematic Diagram of DACRIN Metabolic Model

selected from Pasquill's curves. To provide further flexibility, arbitrarily selected values of standard deviations can be input.

The details of the mathematical models, computer code, data libraries and input preparation are given in Appendices A through F. Sample problem input and output is contained in Appendix G.

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APPENDIX A

Description of Mathematical Models

Reissued October 1975

DESCRIPTION OF MATHEMATICAL MODELS

This appendix describes mathematical models used to calculate inhalation exposures and internal doses from the atmospheric release of radionuclides.

Atmospheric Dispersion Models

The atmospheric dispersion of radionuclides is frequently described mathematically by a bivariate normal distribution model. The standard deviations of the cloud concentration in the crosswind lateral and vertical directions can be estimated from any of the several methods in common use; i.e., Sutton's parameters, Hanford equations, or Pasquill's curves. Using this model, the air concentration at ground level is given by:

$$X = \frac{Q}{\pi \sigma_y \sigma_z \bar{u}_h} \exp \left(-y^2 / 2 \sigma_y^2 - h^2 / 2 \sigma_z^2 \right) \quad (1)$$

where:

- X • ground level air concentration at the coordinates x and y, Ci/m³
- x • downwind distance measured from point of release, meters
- y • crosswind distance measured horizontally from centerline of cloud, meters
- Q • apparent rate of release at receptor, Ci/sec^(a)
- σ_y • crosswind lateral standard deviation of cloud concentration, m
- σ_z • crosswind vertical standard deviation of cloud concentration, m
- \bar{u}_h • average wind speed at the height of release in direction of travel, m/sec
- h • height of release, m

(a) apparent rate of release at receptor is the rate of release at the source corrected for radioactive decay during time of transport to receptor

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Accidental Release

The time-integrated air concentration is convenient to use when calculating doses resulting from accidental releases of short duration. Also, use of the cloud centerline air concentration results in a maximum estimate of this quantity thereby maximizing the resultant dose. The centerline time-integrated air concentration is given by:

$$E_0 = \frac{Q}{\pi \sigma_y \sigma_z \bar{u}_h} \exp(-h^2/2\sigma_z^2) \quad (2)$$

where:

- E_0 • time-integrated air concentration at ground level beneath the centerline of the cloud, Ci·sec/m³
- Q • total release from source, curies.

The crosswind standard deviations, σ_y and σ_z , for stable atmospheric conditions are calculated by the Hanford model^(1,2) as follows:

$$\sigma_y^2 = A [T - \alpha(1 - e^{-T/\alpha})] \quad (3a)$$

$$\sigma_z^2 = a(1 - e^{-k^2 T^2}) + bT \quad (3b)$$

where: T • transport time from point of release to point of interest, sec

and: $A = c + d(\sigma_\theta \bar{u})$

$$\alpha = A/2 (\sigma_\theta \bar{u})^2$$

a , b , c , d , k and $(\sigma_\theta \bar{u})$ are parameters describing the atmospheric condition. Table A-1 summarizes suggested parameter values for use in DACRIN.

TABLE A-1. Values of Meteorological Parameters for the Hanford Model

<u>Parameter</u>	<u>Moderately Stable Conditions</u>	<u>Very Stable Conditions</u>
a	97	34
b	0.33	0.025
c	13	13
d	230	230
k	2.5×10^{-4}	8.8×10^{-4}

MINIMUM VALUES OF $(\sigma_{\theta} \bar{u})$

<u>ASSUMED DURATION OF RELEASE, MIN.</u>	<u>AVERAGE WIND SPEED, m/sec</u>			
	<u>1</u>	<u>2.5</u>	<u>5</u>	<u>10</u>
10	0.024	0.10	0.20	0.30
60	0.04	0.15	0.25	0.35
120	0.08	0.25	0.35	0.45
240	0.10	0.40	0.50	0.60
480	0.18	0.60	0.70	0.90

Crosswind standard deviations calculated for unstable and neutral conditions with Sutton's parameters are:

$$\sigma_y^2 = 0.5C_y^2 x (2 - n) \quad (4a)$$

$$\sigma_z^2 = 0.5C_z^2 x (2 - n) \quad (4b)$$

where:

- n • atmospheric stability index
- C_y • virtual diffusion parameter in the horizontal crosswind direction
- C_z • virtual diffusion parameter in the vertical crosswind direction.

Representative values of the parameters n, C_y and C_z are tabulated in Table A-2.⁽¹⁾

Numerical values of σ_y and σ_z in use at Hanford representative of Pasquill's curves and incorporated in the code are listed in Tables A-3 and A-4.

Inhalation Dose Model

The mathematical model for calculating the dose to an organ of interest via inhalation using the ICRP Task Group Lung Model, TGLM^(3,4) is considerably more complex than that utilized by the lung model initially proposed by the ICRP. In the TGLM, the respiratory tract is divided into three regions, the nasopharyngeal (NP), the tracheobronchial (TB), and the pulmonary (P). The schematic representation of the respiratory tract used in the development of the mathematical model for the deposition and clearance of inhaled radionuclides is shown in Figure 1 of the text. Deposition is assumed to vary with the aerodynamic properties of the aerosol distribution and is described by the three parameters D₃, D₄, and D₅. These parameters represent the fraction of the inhaled material, Q_I initially deposited in the NP, TB and P regions, respectively. Each of the three regions of deposition are further subdivided into two or more subcompartments, each representing the fraction

TABLE A-2. Numerical Values of Atmospheric Dispersion Parameters for Neutral and Unstable Atmospheres

<u>Parameter</u>	<u>Release Level</u>	<u>Wind Speed</u>	<u>Unstable</u>	<u>Neutral</u>
C_y	Ground	1 m/s	0.35	0.21
		5 m/s	0.30	0.15
		10 m/s	0.28	0.14
	Elevated	1 m/s	0.30	0.15
		5 m/s	0.26	0.12
		10 m/s	0.24	0.11
C_z	Ground	1 m/s	0.35	0.17
		5 m/s	0.30	0.14
		10 m/s	0.28	0.13
	Elevated	1 m/s	0.30	0.15
		5 m/s	0.26	0.12
		10 m/s	0.24	0.11
n	—	—	0.20	0.25

TABLE A-3. Values of σ_y for Pasquill Stability Categories

DISTANCE METERS	σ_y for Pasquill Type					
	A	B	C	D	E	F
100	21	16	12	8.0	6.0	3.9
150	34	24	18	12	9.0	6.0
250	54	40	28	20	14	9.8
350	75	55	40	26	20	14
500	100	76	55	37	28	18
700	140	110	76	51	37	26
1,000	200	150	110	72	52	36
1,500	290	220	160	100	75	52
2,500	450	340	240	160	120	81
3,500	610	460	330	220	160	110
5,000	830	630	450	310	220	150
7,000	1,100	840	610	420	300	210
10,000	1,600	1,200	850	570	410	280
15,000	2,200	1,700	1,200	810	570	400
25,000	3,400	2,600	1,800	1,200	880	610
35,000	4,500	3,500	2,500	1,700	1,200	820
50,000	6,200	4,700	3,400	2,300	1,600	1,100
70,000	8,200	6,400	4,700	3,000	2,100	1,500
100,000	11,000	8,500	6,300	4,100	2,800	2,000

TABLE A-4. Values of σ_z for Pasquill Stability Categories

DISTANCE METERS	σ_z for Pasquill Type					
	A	B	C	D	E	F
100	15	10	7.8	4.7	3.0	1.4
150	22	15	11	6.8	4.3	2.2
250	43	26	18	10	7.1	4.0
350	70	37	24	14	9.4	5.3
500	140	57	34	19	13	7.6
700	270	86	46	25	17	10
1,000	670	140	64	33	22	14
1,500	2,000	240	90	43	29	18
2,500	2,000	580	140	62	41	25
3,500	2,000	1,200	190	76	50	30
5,000	2,000	2,000	260	95	61	35
7,000	2,000	2,000	340	120	72	41
10,000	2,000	2,000	440	140	84	47
15,000	2,000	2,000	600	170	99	55
25,000	2,000	2,000	880	220	120	64
35,000	2,000	2,000	1,100	260	130	72
50,000	2,000	2,000	1,400	320	140	79
70,000	2,000	2,000	1,800	370	160	36
100,000	2,000	2,000	2,000	450	170	94

of material initially in a compartment that is subject to a certain clearance process. This fraction is represented by f_j , where j indicates the clearance pathway. The quantity of material in the TB region, for example, cleared by process (c) is then represented by the product $f_c D_4 Q_1$. Values of the (f_j) and the clearance half times T_j for each clearance process for the three translocation classes of aerosols used in the computer code are shown in Table A-5.⁽⁴⁾ Values of the deposition fractions D_3 , D_4 , and D_5 as function of activity median aerodynamic diameter in the form of a graph have been published.⁽³⁾

The respiratory tract model has been incorporated into the simple metabolic model.⁽⁵⁾ Transport of the radionuclides from the respiratory tract, lymphatic systems, and GI tract to other organs and tissues where significant accumulations of the inhaled radionuclide occur, is assumed to take place via the blood. This translocation from the respiratory tract and lymphatic system to the blood has been described in considerable detail.⁽³⁾ Of the material clearing from the respiratory tract through the GI tract, a constant fraction is assumed to be taken up by the blood. Uptake by the n^{th} organ or tissue is assumed to be a constant fraction of the amount entering the blood stream at any time. Once in the n^{th} organ, the activity is assumed to clear the organ and the body at a constant rate.

Notation for the equations found in this section is as follows:

$D_{1n}(T_1)$ is the dose equivalent in rem received by the n^{th} organ or tissue by time T_1 during continuous inhalation of a radioactive aerosol.

$D_{2n}(T_2)$ is the dose equivalent in rem received by the n^{th} organ or tissue by time T_2 following the termination of continuous inhalation of a radioactive aerosol.

$Q_{1n}(T_1)$ is the quantity of radioactive material in μCi present in the n^{th} organ or tissue as a function of time during continuous inhalation of a radioactive aerosol.

TABLE A-5. Values of the Clearance Parameters for the Task Group Lung Model

COMPARTMENT	TRANSLOCATION CLASS						
	j(a)	D		W		Y	
		T_j (b)	f_j (c)	T_j	f_j	T_j	f_j
NP	a	0.01	0.5	0.01	0.1	0.01	0.01
	b	0.01	0.5	0.40	0.9	0.4	0.99
TB	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	e	0.5	0.8	50	0.15	500	0.05
	f	n.a.	n.a.	1	0.4	1	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	50	1	1000	0.9

(a) Metabolic pathways from lung.

(b) Removal half time in days from compartment via pathway j.

(c) Fraction removed from compartment via pathway j.

$Q_{2n}(T_2)$ • is the quantity of radioactive material in μCi present in the nth organ or tissue following the termination of continuous inhalation of a radioactive aerosol.

E_n • is the effective absorbed energy per disintegration in $\frac{\text{MeV}\cdot\text{rem}}{\text{dis}\cdot\text{rad}}$ for the nth organ or tissue.

M_n • is the mass in grams of the nth organ or tissue over which the dose is to be averaged.

λ_j^b • is the biological removal rate constant for the jth sub-compartment of the respiratory tract, sec^{-1} .

λ • is the radiological decay constant of the nuclide of interest, sec^{-1} .

- λ_j • is the total removal rate constant for the k th subcompartment of the respiratory tract, sec^{-1} . Note: $\lambda_j = \lambda_j^b + \lambda$.
- λ_n^b • is the biological removal rate constant for the n th organ or tissue, sec^{-1} .
- λ_n • is the total effective removal rate constant for the n th organ or tissue, sec^{-1} . Note: $\lambda_n = \lambda_n^b + \lambda$.
- f_2 • is the fraction of material in the blood that reaches the organ or tissue of interest.
- f_1 • is the fraction of material in the G.I. tract that reaches the blood.
- P_0 • is the rate at which the radioactive aerosol is inhaled in $\mu\text{Ci}/\text{sec}$.
- f_j • is the fraction of the material in a deposition region, NP, etc., that clears by the j th pathway.
- D_k • is the fraction of the inhaled material deposited in the k th compartment of the respiratory tract.
 $k = 3$ for the nasopharynx (NP) compartment.
 $k = 4$ for the tracheobronchial (TB) compartment.
 $k = 5$ for the pulmonary (P) and lymph (LM) compartments
- T_1 • is the total uptake time, sec.
- T_2 • is the time interval following termination of uptake, sec.

Developing the equations to describe the lung clearance model was divided into two parts. The first part is concerned with describing the organ burdens and the organ doses in the time interval during which the inhalation of radionuclides is taking place. The second part, requiring a different set of equations, describes the organ burdens and the organ doses for the contiguous time interval following the cessation of radionuclide intake.

During uptake, the equations for computing the quantity of radionuclide in the eight subcompartments (j) of the three compartments (k) of the respiratory tract have the form:

$$Q_{j,k}(t) = f_j D_k Q_I \frac{1 - e^{-\lambda_j t}}{\lambda_j} \quad (5)$$

The equations for the removable quantity in the pulmonary lymph nodes are complicated by the fact that both equal and unequal rates for transfer paths into and out of the system are involved. This is due to the dependence of transfer rate upon the solubility class of the radionuclide. Thus, two equations are needed to compute the quantity in the lymph compartment.

For the case $\lambda_h \neq \lambda_i$, i.e., Class Y solubility in the current version of the model:

$$Q_{LMi}(t) = \frac{\lambda_h^b f_i f_h P_0 D_5}{\lambda_h} \left[\frac{(1 - e^{-\lambda_i t})}{\lambda_i} - \frac{(e^{-\lambda_i t} - e^{-\lambda_h t})}{\lambda_h - \lambda_i} \right] \quad (6)$$

and for the case $\lambda_h = \lambda_i$, i.e., Class D and W in the current version of the model:

$$Q_{LMi}(t) = \frac{\lambda_h^b f_i f_h P_0 D_5}{\lambda_h} \left[\frac{(1 - e^{-\lambda_i t})}{\lambda_i} - t e^{-\lambda_i t} \right] \quad (7)$$

With the preceding equations, the equations describing the quantity of radionuclide in the n^{th} organ as a function of time during uptake can be derived by:

$$Q_{In}(t) = P_0 \left\{ \sum_{j=a}^g f_2^j C_j \left[\frac{(1 - e^{-\lambda_n t})}{\lambda_n} - \frac{(e^{-\lambda_n t} - e^{-\lambda_j t})}{(\lambda_j - \lambda_n)} \right] + L \right\} \quad (8)$$

where:

L is the contribution to the organ burden from material passing through the lymphatic system,

and:

$$C_a = \lambda_a^b f_a D_3 / \lambda_a$$

$$C_b = \lambda_b^b f_b D_3 f_1 / \lambda_b$$

$$C_c = \lambda_c^b f_c D_4 / \lambda_c$$

$$C_d = \lambda_d^b f_d D_4 f_1 / \lambda_d$$

$$C_e = \lambda_e^b f_e D_5 / \lambda_e$$

$$C_f = \lambda_f^b f_f D_5 f_1 / \lambda_f$$

$$C_g = \lambda_g^b f_g D_5 f_1 / \lambda_g$$

The lymph pathway contributions to the n^{th} organ burden for the two situations are calculated as follows:

$$L = f_2' C_h \left\{ \frac{1}{\lambda_i} \left[\frac{(1 - e^{-\lambda_n t})}{\lambda_n} - \frac{(e^{-\lambda_n t} - e^{-\lambda_i t})}{(\lambda_i - \lambda_n)} \right] - \frac{1}{(\lambda_h - \lambda_i)} \left[\frac{(e^{-\lambda_n t} - e^{-\lambda_i t})}{(\lambda_i - \lambda_n)} - \frac{(e^{-\lambda_n t} - e^{-\lambda_h t})}{(\lambda_h - \lambda_n)} \right] \right\} \text{ for } \lambda_h \neq \lambda_i \quad (9a)$$

$$L = f_2' C_h \left\{ \frac{1}{\lambda_i} \left[\frac{(1 - e^{-\lambda_n t})}{\lambda_i} - \frac{(e^{-\lambda_n t} - e^{-\lambda_i t})}{(\lambda_i - \lambda_n)} \right] - \frac{e^{-\lambda_n t} - [(\lambda_i - \lambda_n) t + 1] e^{-\lambda_i t}}{(\lambda_i - \lambda_n)^2} \right\} \text{ for } \lambda_h = \lambda_i \quad (9b)$$

$$\text{where: } C_h = \frac{\lambda_i^b \lambda_h^b f_i f_h D_5}{\lambda_h}$$

The calculation of the dose equivalents to organs burdened by the radionuclide are based on the following:

$$D_{1n}(t) = \frac{5.92 \times 10^{-4} E_n}{M_n} \int_0^t Q_{1n}(\tau) d\tau \quad (10)$$

where the constant is a combination of the conversion factors:

$$3.7 \times 10^4 \left(\frac{\text{dis/sec}}{\mu\text{Ci}} \right) 1.6 \times 10^{-6} \left(\frac{\text{ergs}}{\text{MeV}} \right) 10^{-2} \left(\frac{\text{rads}}{\text{erg/g}} \right) = 5.92 \times 10^{-4}$$

The dose to the pulmonary lung at the end of inhalation intake of a radio-nuclide is determined by:

$$D_{lp}(T_1) = \frac{5.92 \times 10^{-4} E_p}{M_p} P_o D_5 \sum_{j=e}^h \frac{f_j}{\lambda_j} (T_1 - A_j) \quad (11)$$

where: M_p is the mass of the lung in grams

$$\text{and: } A_j = \frac{1 - e^{-\lambda_j T_1}}{\lambda_j}$$

For the n^{th} organ or tissue, the dose is computed by:

$$D_{ln}(T_1) = 5.92 \times 10^{-4} \frac{E_n P_o f_2}{M_n} \left\{ \sum_{j=a}^g C_j \left[\frac{T_1 - A_n}{\lambda_n} - \left(\frac{A_n - A_j}{\lambda_j - \lambda_n} \right) \right] \right. \\ \left. + \frac{C_h}{\lambda_i} \left[\frac{T_1 - A_n}{\lambda_n} - \frac{A_n - A_i}{(\lambda_i - \lambda_n)} \right] - \frac{C_h}{(\lambda_h - \lambda_i)} \left[\frac{A_n - A_i}{(\lambda_i - \lambda_n)} \right] \right. \\ \left. - \frac{A_n - A_h}{\lambda_h - \lambda_n} \right\} \text{ for } \lambda_h \neq \lambda_i \quad (12)$$

or:

$$\begin{aligned}
 D_{ln}(T_1) = & 5.92 \times 10^{-4} \frac{E_n P_o f_2'}{M_n} \left(\sum_{j=a}^g C_j \left(\frac{T_1 - A_n}{\lambda_n} - \frac{A_n - A_j}{\lambda_j - \lambda_n} \right) \right. \\
 & + \frac{C_h}{\lambda_i} \left(\frac{T_1 - A_n}{\lambda_n} - \frac{A_n - A_i}{\lambda_i - \lambda_n} \right) + \frac{C_h}{(\lambda_i - \lambda_n)^2} \left\{ A_i - A_n \right. \\
 & \left. \left. + \frac{(\lambda_i - \lambda_n)}{\lambda_i^2} \left[1 - (\lambda_i T_1 + 1) e^{-\lambda_i T_1} \right] \right\} \right) \text{ for } \lambda_h = \lambda_i \quad (13)
 \end{aligned}$$

where:

$$A_n = \frac{1 - e^{-\lambda_n T_1}}{\lambda_n}$$

$$A_i = \frac{1 - e^{-\lambda_i T_1}}{\lambda_i}$$

$$A_h = \frac{1 - e^{-\lambda_h T_1}}{\lambda_h}$$

and C_j and C_h are as previously defined.

Now for the time following cessation of radionuclide inhalation, the equations used to calculate the respiratory tract burden become:

$$Q_{2j}(T_2) = Q_{1j}(T_1) e^{-\lambda_j T_2} \quad (14)$$

where: $Q_{2j}(T_2)$ is the respiratory tract subcompartment burden at a time T_2 following the termination of inhalation uptake.

$Q_{1j}(T_1)$ is the respiratory tract subcompartment burden at the end of inhalation uptake for a time T_1 .

The burden in the n^{th} organ at time T_2 can be described by:

$$\begin{aligned}
 Q_{2n}(T_2) = & Q_{1n}(T_1)e^{-\lambda_n T_2} + f_2' \lambda_i^b Q_{1LMi}(T_1) \frac{(e^{-\lambda_i T_2} - e^{-\lambda_n T_2})}{(\lambda_n - \lambda_i)} \\
 & + f_2' \left\{ \sum_{j=a}^g \frac{C_j'}{(\lambda_j - \lambda_n)} (e^{-\lambda_n T_2} - e^{-\lambda_j T_2}) \right. \\
 & \left. + \frac{C_h'}{(\lambda_h - \lambda_i)} \left[\frac{(e^{-\lambda_n T_2} - e^{-\lambda_i T_2})}{(\lambda_i - \lambda_n)} - \frac{(e^{-\lambda_n T_2} - e^{-\lambda_h T_2})}{(\lambda_h - \lambda_n)} \right] \right\} \text{ for } \lambda_h \neq \lambda_i \quad (15)
 \end{aligned}$$

or by:

$$\begin{aligned}
 Q_{2n}(T_2) = & Q_{1n}(T_1)e^{-\lambda_n T_2} + f_2' \lambda_i^b Q_{1LMi}(T_1) \frac{(e^{-\lambda_i T_2} - e^{-\lambda_n T_2})}{(\lambda_n - \lambda_i)} \\
 & + f_2' \left(\sum_{j=a}^g \frac{C_j'}{(\lambda_j - \lambda_n)} (e^{-\lambda_n T_2} - e^{-\lambda_j T_2}) \right. \\
 & \left. + \frac{C_h'}{(\lambda_i - \lambda_n)} \left\{ e^{-\lambda_n T_2} - [(\lambda_i - \lambda_n)T_2 - 1] e^{-\lambda_i T_2} \right\} \right) \text{ for } \lambda_h = \lambda_i \quad (16)
 \end{aligned}$$

where:

$$C_a' = \lambda_a^b Q_{1a}(T_1)$$

$$C_b' = \lambda_b^b Q_{1b}(T_1) f_1$$

$$C_c' = \lambda_c^b Q_{1c}(T_1)$$

$$C_d' = \lambda_d^b Q_{1d}(T_1) f_1$$

$$C_e' = \lambda_e^b Q_{1e}(T_1)$$

$$C_f' = \lambda_f^b Q_{1f}(T_1) f_1$$

$$C_g' = \lambda_g^b Q_{1g}(T_1) f_1$$

$$C_h' = f_i \lambda_h^b \lambda_i^b Q_{1h}(T_1)$$

The pulmonary lung dose from inhalation of a radionuclide for a time T_1 followed by no additional radioactive intake for a time T_2 is determined by:

$$D_p = D_{1p}(T_1) + 5.92 \times 10^{-4} \frac{E_p P_0 D_5}{M_p} \sum_{j=e}^h f_j A_j B_j \quad (17)$$

$$\text{where: } B_j = \frac{1 - e^{-\lambda_j T_2}}{\lambda_j}$$

and $D_{1p}(T_1)$ and A_j are the same as previously defined by Equation (11).

The effective dose to the n^{th} organ as a result of inhalation uptake for a time T_1 followed by no additional radioactive intake for a time T_2 is determined by:

$$D_n = D_{1n}(T_1) + 5.92 \times 10^{-4} \frac{E_n}{M_n} \left\{ Q_{1n}(T_1) B_n + f_2 \left[\frac{\lambda_i^b Q_{1LMi}(T_1)}{(\lambda_n - \lambda_i)} (B_i - B_n) \right. \right. \\ \left. \left. + G \left[\frac{(B_n - B_i)}{(\lambda_i - \lambda_n)} - Z \right] + \sum_{j=a}^g \frac{C_j'}{(\lambda_j - \lambda_n)} (B_n - B_j) \right] \right\} \quad (18)$$

$$\text{where: } B_n = \frac{1 - e^{-\lambda_n T_2}}{\lambda_n}$$

$$B_h = \frac{1 - e^{-\lambda_h T_2}}{\lambda_h}$$

$$B_i = \frac{1 - e^{-\lambda_i T_2}}{\lambda_i}$$

$$B_j = \frac{1 - e^{-\lambda_j T_2}}{\lambda_j}$$

also, for $\lambda_h \neq \lambda_i$:

$D_{1n}(T_1)$ is determined by Equation (12)

$Q_{1n}(T_1)$ is determined by Equations (8,9a)

$$G = \frac{C'_h}{(\lambda_h - \lambda_i)}$$

$$Z = \frac{B_n - B_h}{(\lambda_h - \lambda_n)}$$

and for $\lambda_h = \lambda_i$:

$D_{1n}(T_1)$ is determined by Equation (13)

$Q_{1n}(T_1)$ is determined by Equations (8,9b)

$$G = \frac{C'_h}{(\lambda_i - \lambda_n)}$$

$$Z = \lambda_i^{-2} [1 - (\lambda_i T_2 + 1) e^{-\lambda_i T_2}]$$

Provision is made in the code to calculate the dose equivalent to the pulmonary lung as a result of primary deposition, i.e., that occurring directly on the lung surfaces during inhalation and also for secondary deposition, i.e., that occurring as a result of solubilization and redeposition in the lung tissue itself, in this respect the pulmonary lung is treated as any other organ.

Weighted values of the effective energy of daughter radionuclides, ϵ_i were calculated using the ICRP equations⁽⁶⁾ for three biological clearance half times of 1 day, 50 days and 500 days. The effective energy of the i^{th} daughter is determined by:

$$\epsilon_i = \sum_j F_j [E(\text{RBE})_j]_i$$

where:

$$F_i = \frac{\prod_{j=1}^i T_j / T_j^r}{1 - e^{-\lambda_0 t}} \sum_{n=1}^i \frac{T_n^i (1 - e^{-\lambda_n t})}{\prod_{\substack{p=0 \\ p \neq n}}^i (T_n - T_p)}$$

and:

- F_i • the ratio of the number of disintegrations of daughter atoms to the number of disintegrations of parent atoms in the lung over a time t .
- λ_0 • the effective decay constant of the parent, the subscript, zero, refers to the parent.
- λ_i • the effective decay constant of the i^{th} daughter in the lung.
- T_i • the effective half life of the i^{th} daughter in the lung.
- T_i^r • the radioactive half life of the i^{th} daughter.
- t • the time of interest over which the dose is to be calculated.
- E • the total energy absorbed in the organ per disintegration of the radionuclide.
- RBE • the relative biological effectiveness of the radiation.
- n • the relative damage factor for radionuclides deposited in the bone.

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2. J. J. Fuquay, C. L. Simpson and W. T. Hinds, Estimates of Ground Level Air Exposures Resulting from Protracted Emissions from 70-Meter Stacks at Hanford, HW-80204, 1964.
3. Task Group of Committee 2, ICRP, Task Group on Lung Dynamics for Committee II of the ICRP, Health Physics, Vol. 12, p. 173, 1966.
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5. Paul G. Volilleque AERIN, A Code for the Acute Aerosol Inhalation Exposure Calculations, Health Physics, Vol. 19, p. 427, 1970.
6. Report of Committee II on Permissible Dose for Internal Radiation, ICRP Publication 2, Pergamon Press, 1959.

APPENDIX B

COMPUTER CODE

APPENDIX B

COMPUTER CODE

The models described in Appendix A have been incorporated into a computer code programmed for the CDC CYBER 74 Computer.

MAIN

MAIN is the main program and calls the other routines as needed to perform the calculations requested by the user. The data library reading subroutines are called by the main program and all other input data is read in this program. Input data that is required is explained in detail in Appendix F. If the calculation of E/Q values is called for, this program calls any required subroutines to assist. Maximum flexibility is provided in the generation of E/Q values. The code allows the use of Hanford Model equations, Sutton Model equations, data taken from Pasquill curves for σ_z and σ_y , the input of σ_y and σ_z of the users choice or the input of E/Q values directly. If the activity median aerodynamic diameter is input for the aerosol, the routine to generate the deposition fractions is called. If fission products input has been specified, the routine to perform a chain decay for the time during which the nuclides are in transit from the release point to the receptor is called. The ORDOSE subroutine is then called to calculate the doses to the organs chosen.

ORDOSE

Subroutine ORDOSE controls the routines that generate the necessary values for the organ dose calculation. All printing of the individual nuclide contributions to the organ dose values as well as the totals is accomplished in this routine. All values returned by called subroutines, ORDOT1, ORDOT2, QLUNT1, QLMIT1, and QNT1, are normalized to a unit uptake rate. These values are then multiplied by the actual uptake rate to arrive at the actual dose to the organ. Output page format is determined in this routine by the number of organs chosen at execution time (up to a maximum

of 10). Where no data exists for a nuclide-organ combination, a blank shows on the output page for that combination rather than a zero.

DEP

Subroutine DEP is used to calculate the compartment deposition fractions D_3 , D_4 and D_5 when the activity median aerodynamic diameter is input instead of the fractions. The error function is integrated numerically over appropriate limits to produce the necessary fractions.

SIMPS

Function SIMPS performs the numerical integration of the error function using Simpsons rule.

ERFXP

Function ERFXP is the error function: e^{-x^2} that is integrated by SIMPS.

PASSIG

Subroutine PASSIG interpolates on a table of stored Pasquill data values of σ_y and σ_z at fixed distances and produces the values of Pasquill σ_y and σ_z for the distances of interest. The σ_y and σ_z data is in the form of data statements for each Pasquill atmospheric stability class A through F at 20 distances from release point out to 10^7 meters.

UNIT

Subroutine Unit converts input time in seconds to more convenient units depending on the magnitude of the value input. The appropriate units, i.e., minutes, hours, days, etc., are also returned for use in the printed output.

CHAINT

Subroutine CHAINT performs a chain decay on all fission product decay chains from which nuclides for consideration have been chosen. The time over which the decay is performed is the source to receptor transit time.

ORDOT1

Function ORDOT1 calculates the dose to an organ for the time T_1 during which inhalation of activity occurs at a unit rate.

ORDOT2

Function ORDOT2 calculates the dose to an organ during a time T_2 following inhalation of activity at a unit rate. The equations used, take into account any continuing movement of material from the respiratory compartments and the lymph to the organ following the termination of the inhalation of radioactive material.

QLUNT1

Function QLUNT1 calculates the activity in each lung subcompartment following continuous inhalation at a unit rate for a time T_1 .

QLMIT1

Function QLMIT1 calculates the quantity of a nuclide present in the pulmonary lymph, that is removable, following continuous uptake at a unit rate for a time T_1 .

QNT1

Function QNT1 calculates the quantity of a nuclide present in an organ following continuous uptake at a unit rate for a time T_1 .

The results from the preceding three routines are used in the calculation of the dose following cessation of inhalation uptake.

LIBB

Subroutine LIBB reads the biological data from the library and converts some of the values into units more readily used by the other routines in the program.

LIBK

Subroutine LIBK reads the nuclide radiological data library, converts some of the values to other units and generates the fractional decay yield array for use in the fission product chain calculations.

APPENDIX C

CODE FLOW DIAGRAM

APPENDIX C

CODE FLOW DIAGRAM

A general outline of the code operation is included here. The relationship between the main program and subroutines used by the code is shown in Figure C-1. The code flow diagram shown in Figure C-2 through C-12 covers the logic in the main program and subroutine ORDOSE in some detail. Code flow diagrams for the auxiliary routines are not shown, however, their function is included in the blocks at the appropriate places within the flow diagrams.

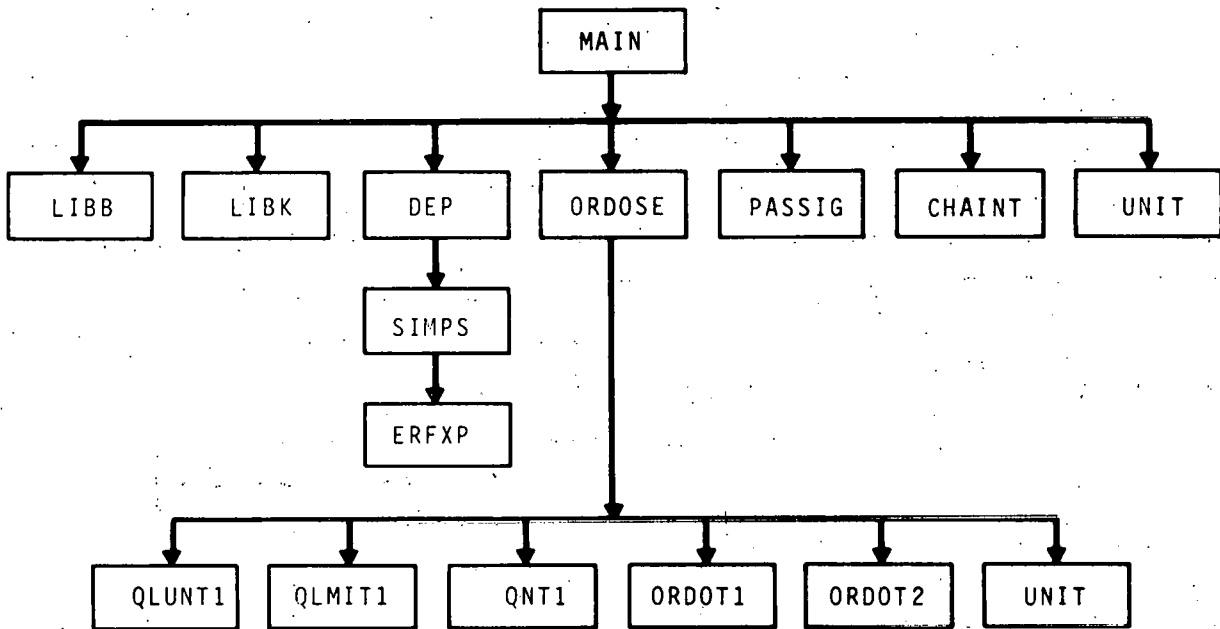


FIGURE C.12. Code Flow Diagram

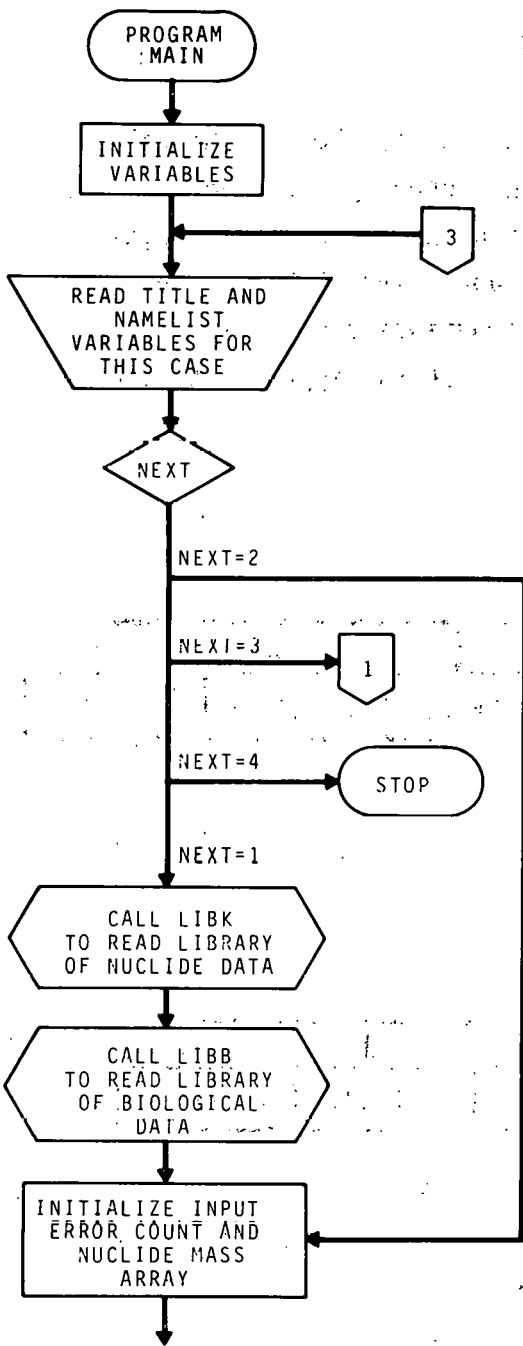


FIGURE C.2. Code Flow Diagram

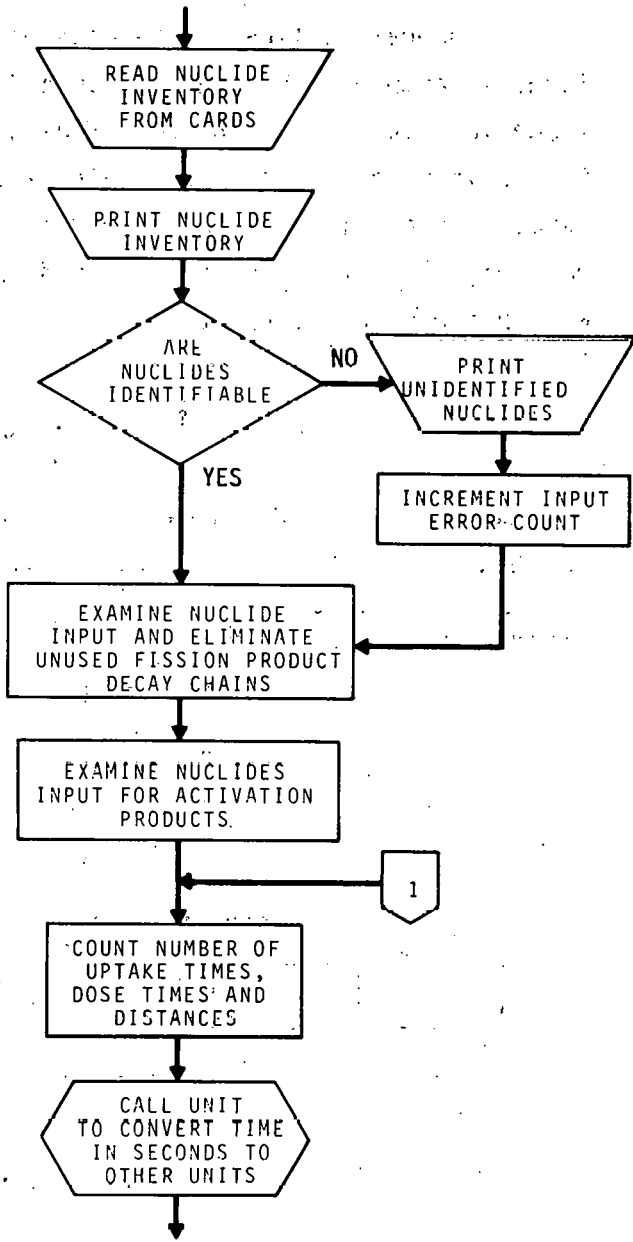


FIGURE C.3. Code Flow Diagram

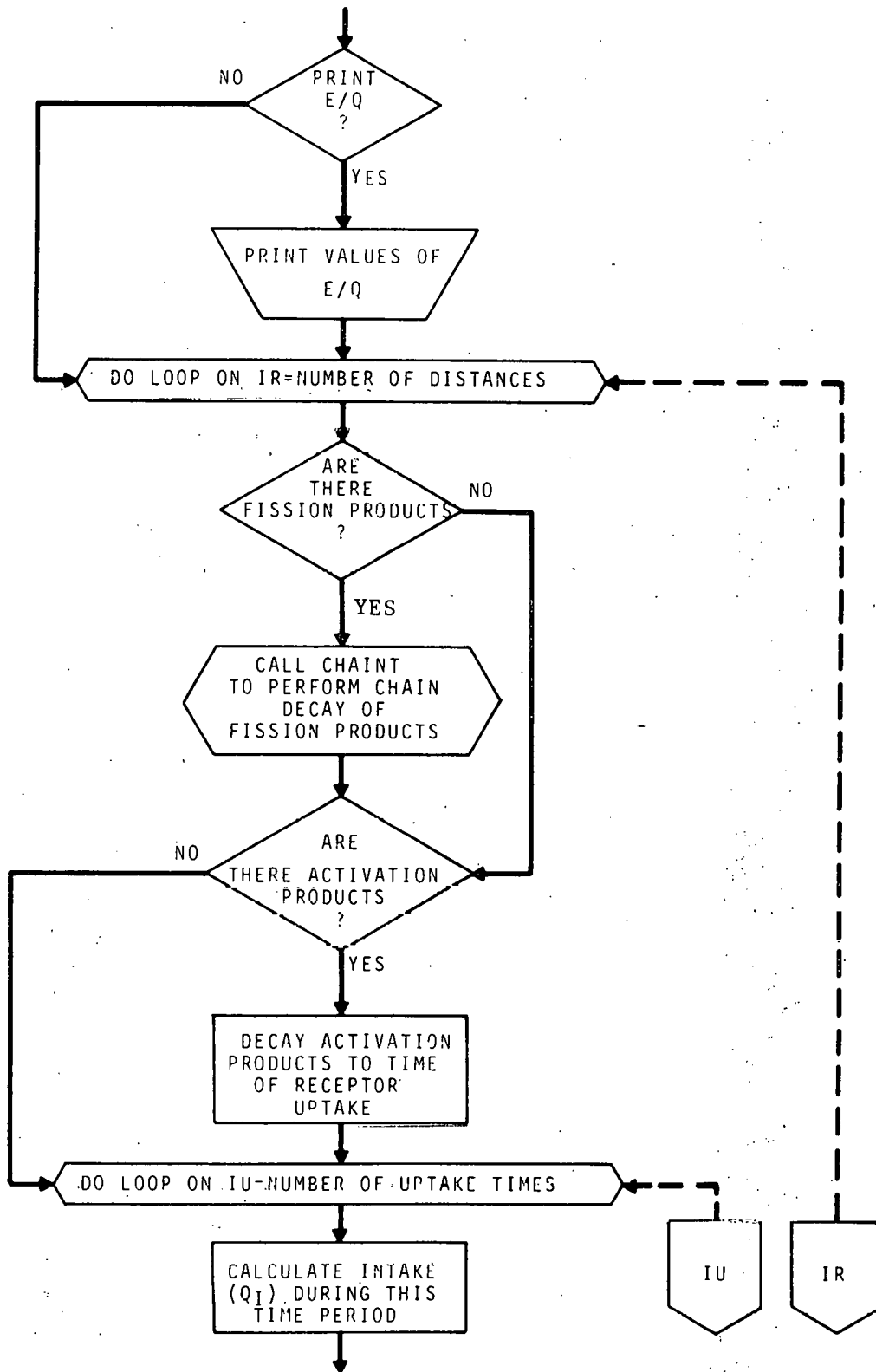


FIGURE C.5. Code Flow Diagram

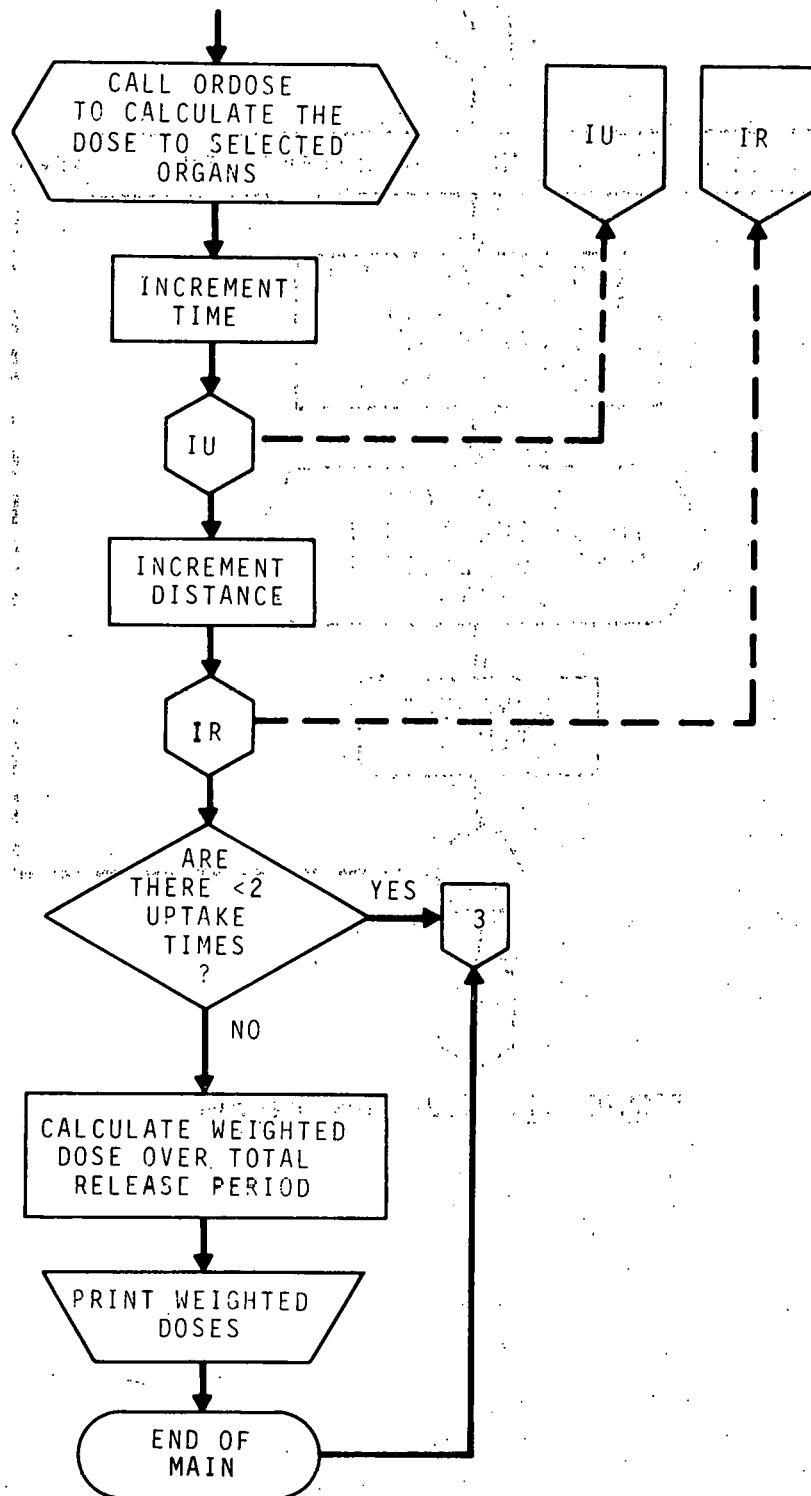


FIGURE C.6. Code Flow Diagram

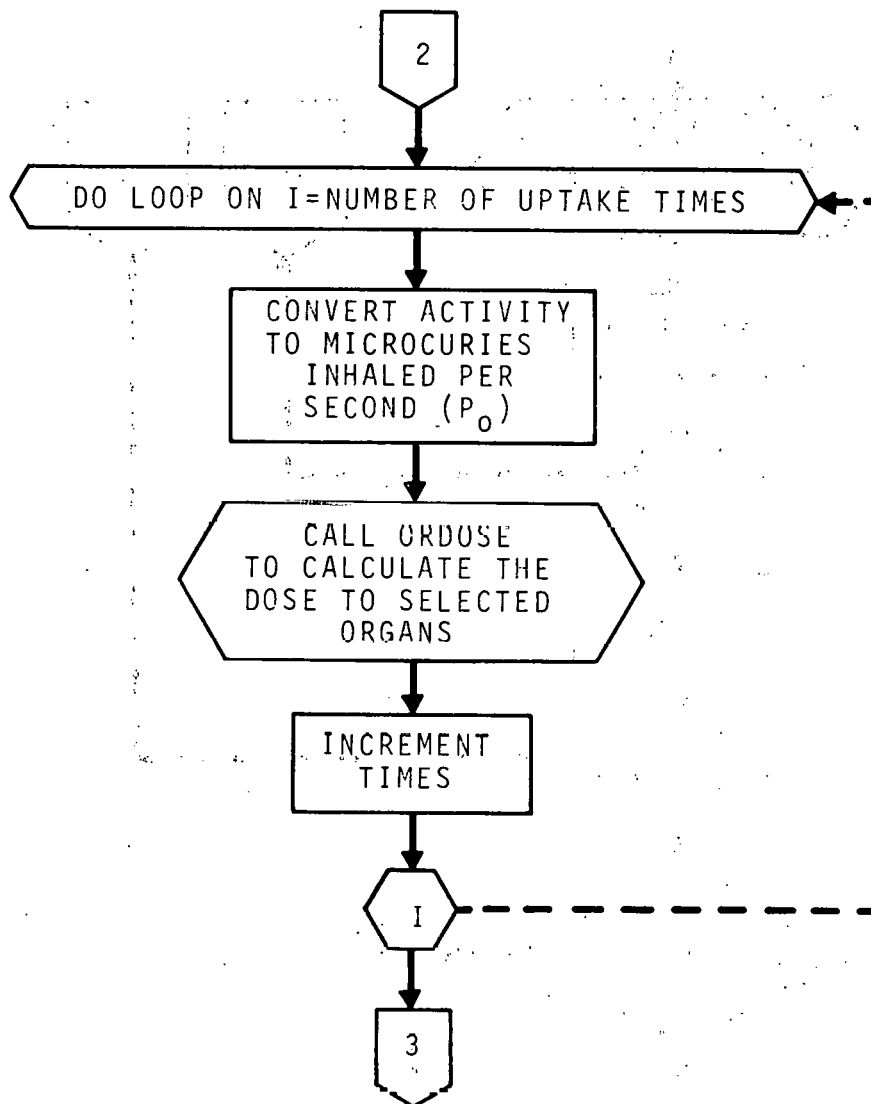


FIGURE C.7. Code Flow Diagram

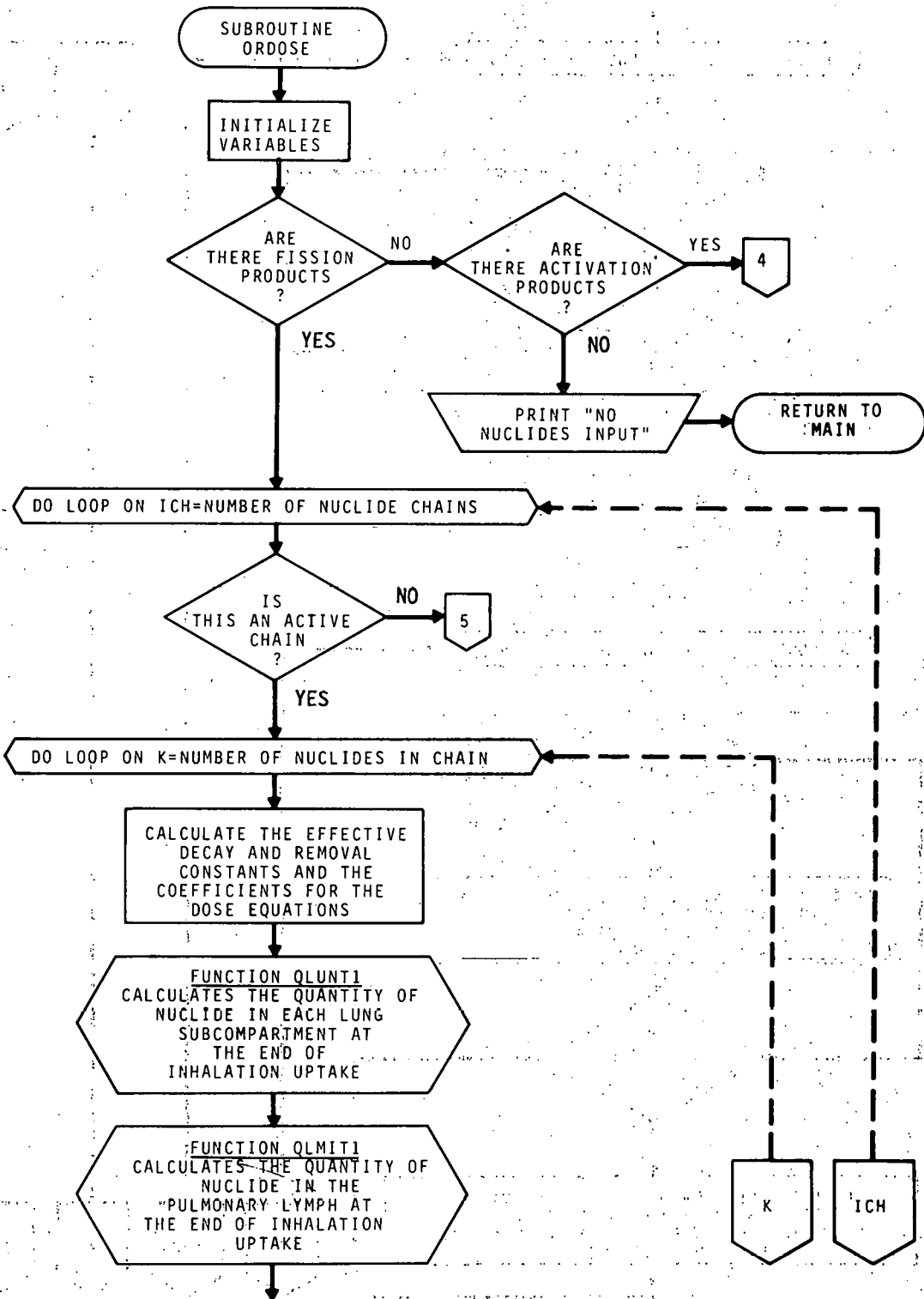


FIGURE C.8. Code Flow Diagram

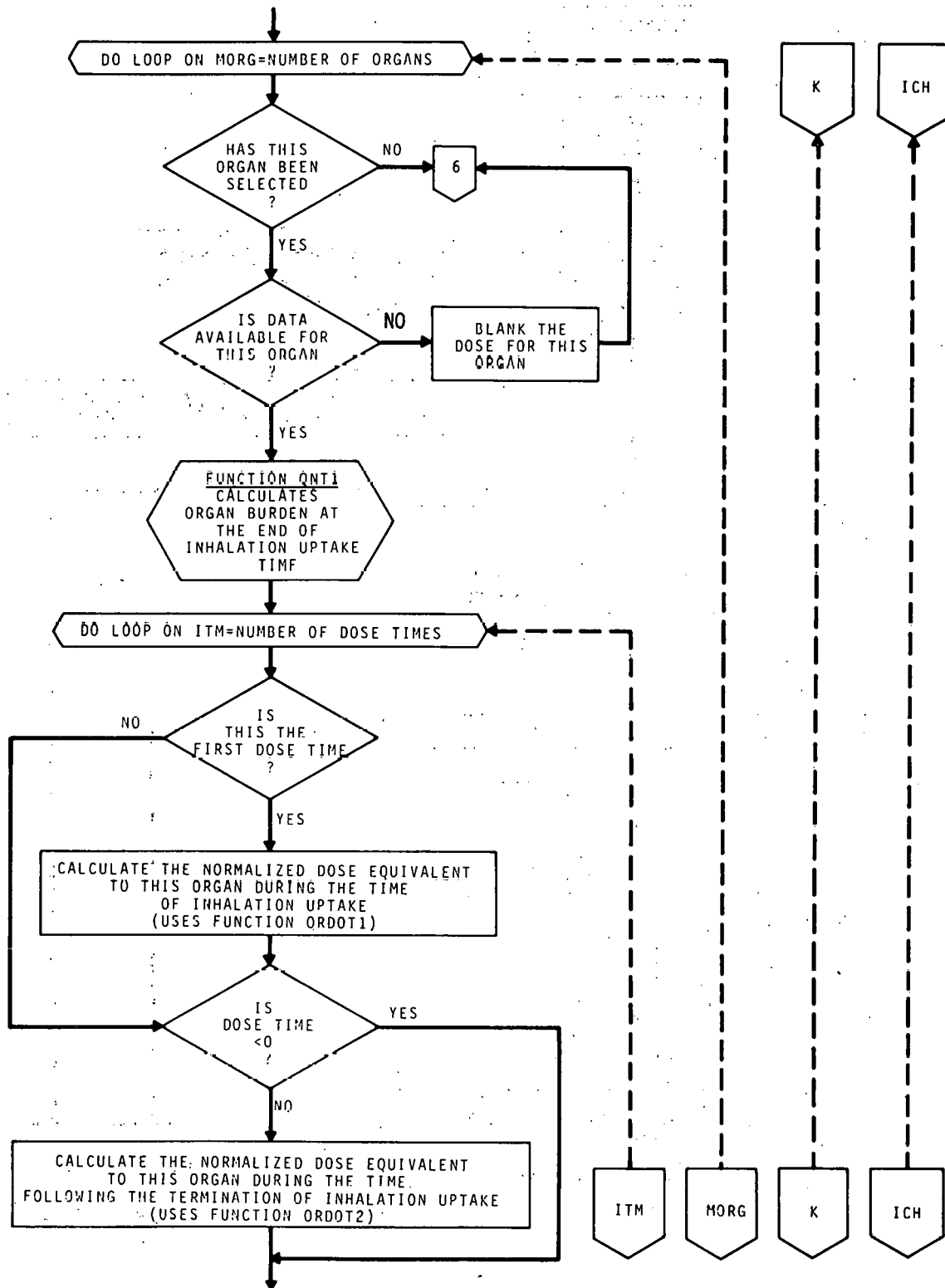


FIGURE C.9. Code Flow Diagram

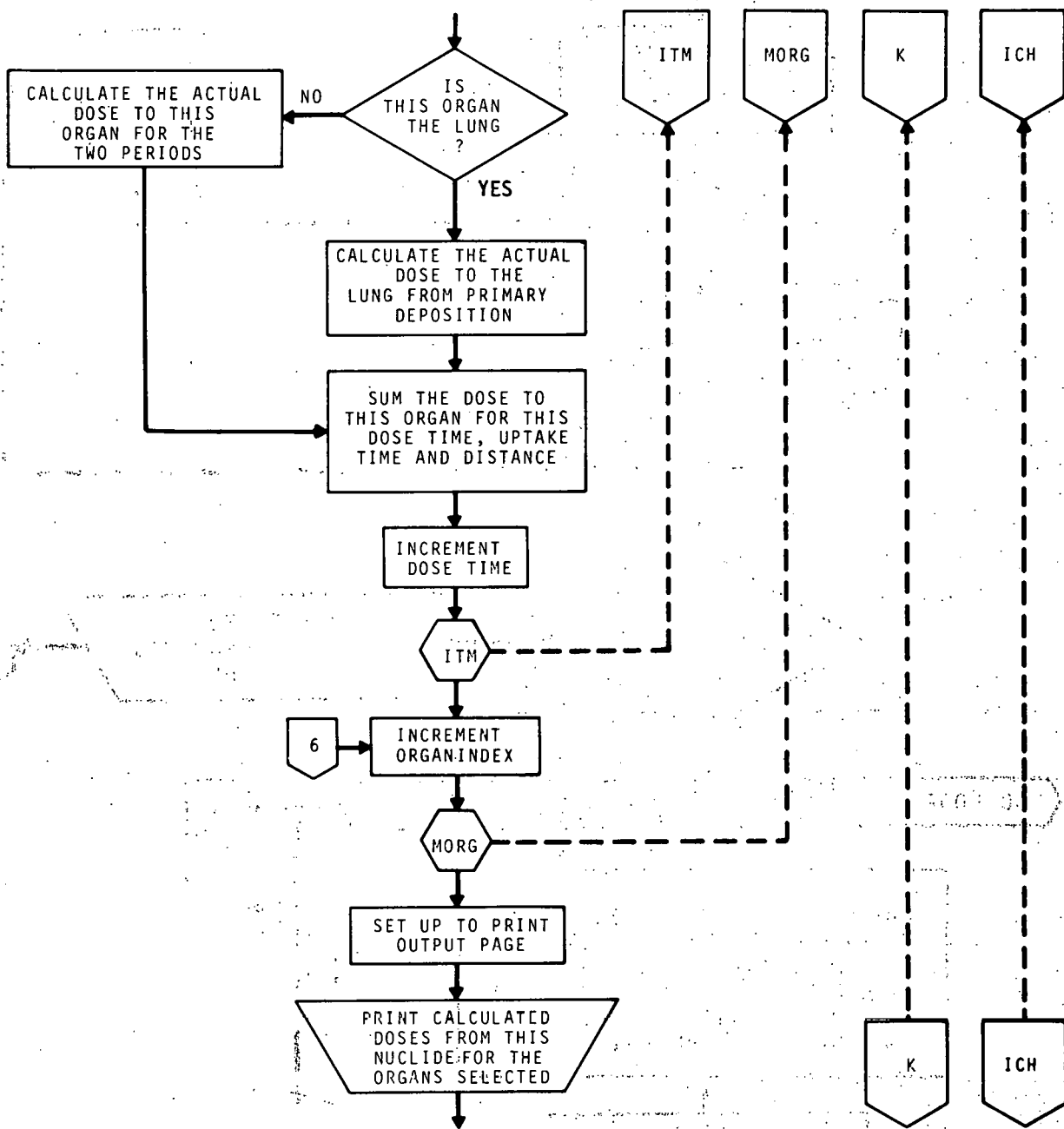


FIGURE C.10. Code Flow Diagram

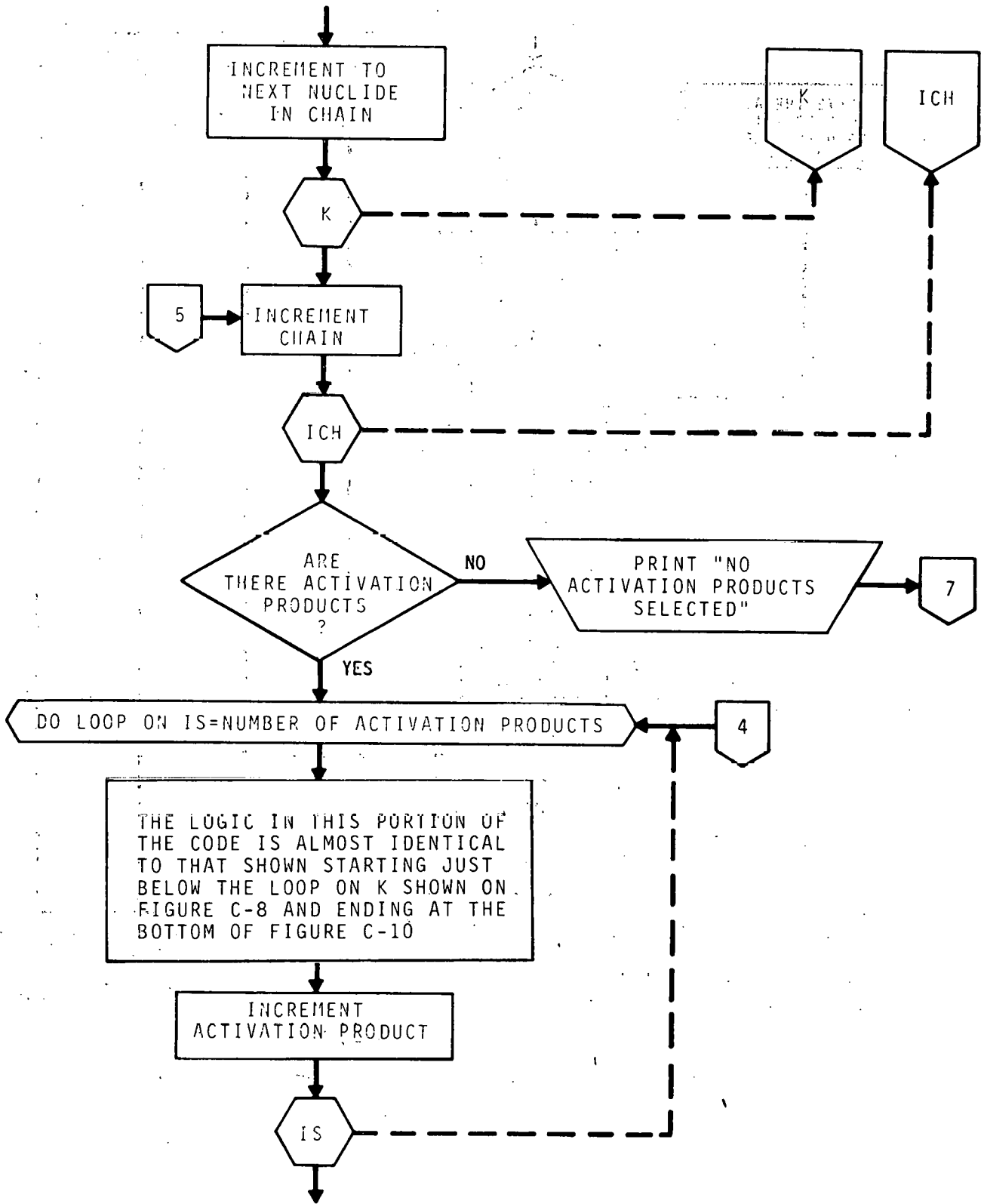


FIGURE C.11. Code Flow Diagram

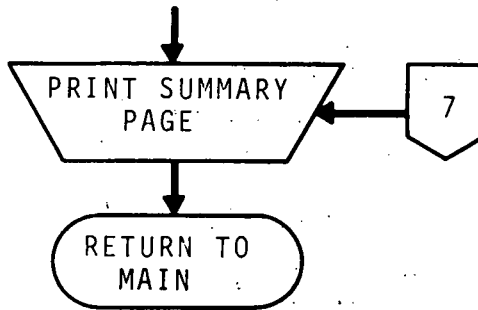


FIGURE C.12. Code Flow Diagram

APPENDIX D

DATA LIBRARIES

APPENDIX D

DATA LIBRARIES

Two data libraries are used by the code.

1. Isotope Data Library
2. Organ Data Library

The data in the libraries is set up such that additional data or changes to existing data may be made with little difficulty.

ISOTOPE DATA LIBRARY

The data in this library is used by CHAINT in performing the chain decay of fission products specified in the input data. Half-life data for activation product nuclides is included in the second half of this library. The library has been taken from that originally set up for a reactor fission product inventory code called RIBD,⁽¹⁾ hence not all of the data is required by the DACRIN code. The RIBD code could be added at a later date to generate a fission product inventory for input to this code.

The first card image of the isotope library contains the number of fission products in the library in I3 format in the first three columns. There is one card image for each fission product arranged in ascending atomic number order within an ascending atomic mass sequence. Currently the library contains data on 450 fission products.

The card image after the last fission product card image contains the number of activation products in I3 format in the first three columns. There is one card image for each activation product nuclide and there are currently 144 activation product nuclides in the library.

<u>COLUMN</u>	<u>FORMAT</u>	<u>VARIABLE</u>
1-3	I3	Atomic Mass Number
4-5	I2	Atomic Number
6-14	E9.3	Physical half-life, days
15-23	E9.3	*Fission yield from species one (^{235}U), %
24-32	E9.3	*Fission yield from species two (^{239}Pu), %
33-41	E9.3	*Absorption cross section, barns
42-46	E5.3	*Fraction of (n, α) captures that go to isomeric state
47-51	E5.3	If in isomeric state, fraction of decays by beta emission; if in ground state, fraction that decays to an isomer
52-56	E5.3	*Average beta energy released per disintegration, MeV
57-61	E5.3	*Average gamma energy released per disintegration, MeV
69-71	A3	Isotope name (left adjusted)
72	I1	*Isotope type: 1-Isomeric state 2-Ground state

*Not used by code.

ORGAN DATA LIBRARY

This library contains most of the data required by ORDOSE to calculate the effective dose to an organ. Data in the library is based on recommendations of the ICRP. (2,3,4)

The first card image contains the total number of data card images in the library in I4 format in the first four columns. The present limit is 1200 card images. There is one card image for each radionuclide-organ combination, (where data exists), arranged in ascending organ number within an ascending atomic mass sequence within an ascending atomic number order. There are currently 1136 entries in this library.

<u>COLUMN</u>	<u>FORMAT</u>	<u>VARIABLE</u>
1-3	I3	Atomic Number
4-6	I3	Atomic Mass Number
7	I1	Isotope type: blank - ground state 1 - Isomeric state
8-9	A2	Element name
10-11	I2	Organ code number
12-19	E8.3	Physical half-life, days
21-27	E8.3	Biological half-life in the organ or tissue, days
28-35	E8.3	*Fraction of that material taken into the body by ingestion that is retained in the organ, f_w in the initial ICRP lung model.
36-43	E8.3	**Fraction of that taken into the body by inhalation that arrives in the organ, f_a in the initial ICRP lung model.
44-51	E8.3	The fraction of the nuclide in the blood reaching the organ, f_2
52-59	E8.3	Effective absorbed energy per disintegration of the radionuclide and daughters in the organ, E, MeV (in the case of the lung this is for a 1 day half residence time)
60-65	E6.2	Weighted value of the effective absorbed energy per disintegration of the radionuclide and its daughters in the lung for a 50 day half residence time
66-71	E6.2	Same as previous field except for a 500 day half residence time in the lung.

* Used for extracting f_1 , the fraction ingested reaching the blood.
 ** Not used by code.

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3. Recommendations of the International Commission on Radiological Protection, ICRP Publication 6, Pergamon Press, 1962.
4. The Metabolism of Compounds of Plutonium and Other Actinides, A Report prepared by a Task Group of Committee 2 of the International Commission on Radiological Protection, ICRP Publication 19, Pergamon Press, May 1972.

ISOTOPE DATA LIBRARY

450

72301.938E-01	1.600E-05	1.200E-04	0.	0.	.12	1.235	ZN 2	RIBLIB	2
72315.875E-01	0.	0.	0.	0.	.43	2.869	GA 2	RIBLIB	3
7232	0.	0.	1.	0.			GE 2	RIBLIB	4
73301.390E-03	9.800E-05				2.25		ZN 2	RIBLIB	5
73312.000E-01	2.000E-06				.42	.409	GA 2	RIBLIB	6
73320	0.	0.	14.	0.			GE 2	RIBLIB	7
74315.555E-03	3.500E-04				1.01	3.149	GA 2	RIBLIB	8
7432	0.	0.	.5	.4			GE 2	RIBLIB	9
75311.389E-03	4.000E-04				1.43	.58	GA 2	RIBLIB	10
75325.671E-04	4.000E-04					.139	GE*1	RIBLIB	11
75325.694E-02					.42	.938	GE 2	RIBLIB	12
7533	0.	0.	4.5	0.			AS 2	RIBLIB	13
76313.704E-04	.001				2.74	1.	GA 2	RIBLIB	14
7632	.001		.2	.5			GE 2	RIBLIB	15
76331.104E-00	0.	0.			1.09	0.404	AS 2	RIBLIB	16
7634	0.	0.	85.	.259			SE 2	RIBLIB	17
77326.250E-04	5.400E-03				0.640	.77	GE*1	RIBLIB	18
77324.583E-01	3.700E-03				.64	1.134	GE 2	RIBLIB	19
77331.625E-00	0.				.22	0.022	AS 2	RIBLIB	20
77342.083E-04						.16	SE*1	RIBLIB	21
7734	0.	0.	42.	0.			SE 2	RIBLIB	22
78328.750E-02	1.800E-02				.32	0.300	GE 2	RIBLIB	23
78336.319E-03	2.000E-03				1.47	0.888	AS 2	RIBLIB	24
7834	0.	0.	.41	.875			SE 2	RIBLIB	25
79336.250E-03	4.000E-02				1.	.90	AS 2	RIBLIB	26
79342.708E-03	0.					0.096	SE*1	RIBLIB	27
79342.555E-07	0.				.04		SE 2	RIBLIB	28
79350	0.	0.	1.140E	0.10.2540.			BR 2	RIBLIB	29
80331.736E-04	.08				2.73	.5	AS 2	RIBLIB	30
8034			.6	.2			SE 2	RIBLIB	31
80351.875E-01	0.					0.085	BR*1	RIBLIB	32
80351.250E-02	0.				.72	.116	BR 2	RIBLIB	33
8036	0.	0.	15.	0.			KR 2	RIBLIB	34
81333.819E-04	1.250E-01				1.66		AS 2	RIBLIB	35
81343.958E-02	8.400E-03					0.103	SE*1	RIBLIB	36
81341.250E-02	0.				.53		SE 2	RIBLIB	37
8135	0.	0.	3.	0.			BR 2	RIBLIB	38
8136	0.	0.					KR 2	RIBLIB	39
8234	3.800E-05		5.400E-02	0.9200.			SE 2	RIBLIB	40
82351.471E-00					.14	2.648	BR 2	RIBLIB	41
8236	0.	0.	45.	1.			KR 2	RIBLIB	42
83347.986E-04	3.000E-01	1.200E-01			1.000	1.55	SE*1	RIBLIB	43
83341.736E-02	1.800E-01	8.600E-02			.69	1.000	SE 2	RIBLIB	44
83351.000E-01	8.400E-02	0.	0.	1.000	.34	0.010	BR 2	RIBLIB	45
83367.750E-02	0.	0.	0.	0.	0.	0.041	KR*1	RIBLIB	46
8336	0.	0.	180.	0.			KR 2	RIBLIB	47
84342.083E-03	6.100E-02	3.000E-02			.49		SE 2	RIBLIB	48
84354.167E-03	1.900E-02	1.000E-02			1.	.71	BR*1	RIBLIB	49
84352.222E-02	9.200E-01	4.300E-01			1.23	1.863	BR 2	RIBLIB	50
8436	0.	0.	.14	.714			KR 2	RIBLIB	51
85334.977E-06	5.500E-01	1.1.260E-01			1.28	5.0	AS 2	RIBLIB	52
85344.514E-04	5.500E-01	1.1.260E-01			2.24		SE 2	RIBLIB	53
85352.083E-03	4.000E-01	1.1.600E-01			1.	1.04	BR 2	RIBLIB	54
85361.833E-01	0.	0.			0.810	.29	KR*1	RIBLIB	55
85363.927E-03	1.270E-01	1.15.			.25	0.004	KR 2	RIBLIB	56
8537	0.	0.	1.000E	0.00.1			RB 2	RIBLIB	57
86341.952E-04	1.010E-03	3.800E-01			1.76		SE 2	RIBLIB	58
86356.250E-04	1.010E-03	3.800E-01			3.23	.9	BR 2	RIBLIB	59
8636			.06				KR 2	RIBLIB	60
86377.222E-04	0.	0.				0.56	RB*1	RIBLIB	61
86371.870E-01	2.800E-05	2.300E-05			.66	0.097	RB 2	RIBLIB	62
8638	0.	0.	.006	1.			SR 2	RIBLIB	63
87341.852E-04	2.700E-09	2.00E-01			3.73		SE 2	RIBLIB	64
									65

87356.366E-040.	0.		1.97	3.780	BR 2	RIBLIB	66
87365.273E-020.	0.	630.	1.34	1.089	KR 2	RIBLIB	67
8737	0.	0.	.12		RB 2	RIBLIB	68
87381.167E-010.	0.			0.388	SR*1	RIBLIB	69
8738	0.	0.			SR 2	RIBLIB	70
88351.852E-042.900E 001.100E 00			3.22	.035	BR 2	RIBLIB	71
88361.167E-018.000E-013.200E-01			.38	1.922	KR 2	RIBLIB	72
88371.250E-020.	0.	1.	1.99	.748	RB 2	RIBLIB	73
8838		.005			SR 2	RIBLIB	74
89355.208E-052.000E-017.000E-02			3.	.075	BR 2	RIBLIB	75
89362.222E-034.400E 001.640E 00			1.39	2.265	KR 2	RIBLIB	76
89371.042E-022.000E-010.	.55		.60	2.455	RB 2	RIBLIB	77
89385.04 E 010.	0.	.4	.0002	.58	SR 2	RIBLIB	78
89391.852E-04				.915	Y* 1	RIBLIB	79
8939	0.	0.	1.301	.00080.	Y 2	RIBLIB	80
90351.852E-052.600E 009.800E-01			3.	.075	BR 2	RIBLIB	81
90363.819E-042.600E 009.800E-01			1.37	2.23	KR 2	RIBLIB	82
90372.014E-037.000E-012.900E-01			2.05	4.15	RB 2	RIBLIB	83
90381.022E 040.	0.	1.	.20		SR 2	RIBLIB	84
90391.333E-01				.68	Y* 1	RIBLIB	85
90392.675E 000.	0.		.94	.0003	Y 2	RIBLIB	86
9040	0.	0.	.1		ZR 2	RIBLIB	87
91361.157E-043.700E 001.260E 00			.5	1.56 2.	KR 2	RIBLIB	88
91379.722E-032.000E-017.000E-02			1.	1.27 2.8	RB*1	RIBLIB	89
91378.338E-042.000E 001.100E 00				2.04 0.4	RB 2	RIBLIB	90
91384.042E-010.	0.		.590	.63 0.742	SR 2	RIBLIB	91
91393.472E-020.	0.			0.551	Y* 1	RIBLIB	92
91395.900E 010.	0.	1.4	.61	0.004	Y 2	RIBLIB	93
9140	0.	0.	1.		ZR 2	RIBLIB	94
92363.471E-052.700E 001.530E 00				2.73	KR 2	RIBLIB	95
92375.787E-052.800E 001.540E 00				2.97	RB 2	RIBLIB	96
92381.125E-016.000E-017.000E-02				2.26 1.274	SR 2	RIBLIB	97
92391.471E-010.	0.			1.44 0.266	Y 2	RIBLIB	98
9240	0.	0.	.2		ZR 2	RIBLIB	99
93362.315E-051.300E 007.700E-01				3.82	KR 2	RIBLIB	100
93376.944E-053.100E 001.820E 00				2.83	RB 2	RIBLIB	101
93385.764E-032.000E 001.320E 00				1.40 1.0	SR 2	RIBLIB	102
93394.208E-011.000E-016.000E-02				1.18 0.102	Y 2	RIBLIB	103
93403.468E 080.	0.	4.	.25	.015	ZR 2	RIBLIB	104
93411.351E 030.	0.			0.029	NR*1	RIBLIB	105
9341	0.	0.	1.1	.909	NR 2	RIBLIB	106
94361.620E-056.000E-016.000E-01				2.70	KR 2	RIBLIB	107
94373.472E-052.300E 001.500E 00				4.57	RB 2	RIBLIB	108
94388.333E-042.900E 001.640E 00				1.46	SR 2	RIBLIB	109
94391.389E-027.000E-017.400E-01				2.23 1.	Y 2	RIBLIB	110
9440	0.	0.	.08		ZR 2	RIBLIB	111
94414.583E-030.	0.		.001	.00050.042	NR*1	RIBLIB	112
94417.300E 060.	0.	15.		.16 1.570	NR 2	RIBLIB	113
9442	0.	0.			MO 2	RIBLIB	114
95361.157E-053.200E 002.900E 00				4.57	KR 2	RIBLIB	115
95372.314E-053.200E 002.900E 00				3.58	RB 2	RIBLIB	116
95385.556E-04				2.72	SR 2	RIBLIB	117
95397.292E-03				2.09	Y 2	RIBLIB	118
95406.500E 010.	0.		.02	.12 0.720	ZR 2	RIBLIB	119
95413.750E 000.	0.			0.235	NR*1	RIBLIB	120
95413.500E 010.	0.	7.	.043	0.760	NR 2	RIBLIB	121
9542	0.	0.	14.		MO 2	RIBLIB	122
96391.597E-033.200E 002.585E 00				1.55 1.7	Y 2	RIBLIB	123
9640		3.200E 002.585E 00.05			ZR 2	RIBLIB	124
96419.583E-015.700E-043.600E-03				.22 2.456	NR 2	RIBLIB	125
9642	0.	0.	1.		MO 2	RIBLIB	126
97361.157E-052.400E 002.210E 00				4.20	KR 2	RIBLIB	127
97372.315E-052.400E 002.210E 00				4.39	RB 2	RIBLIB	128
97383.472E-05				3.57	SR 2	RIBLIB	129
97394.630E-05				2.31	Y 2	RIBLIB	130
97407.983E-011.400E 001.230E 00			.9	.70 0.221	ZR 2	RIBLIB	131

97416.944E-040.	0.			0.75	NB*1 RIBLIB	132
97415.000E-020.	0.			.47 .673	N9 2 RIBLIB	133
9742	0.	2.			MO 2 RIBLIB	134
98406.944E-04 .2	1.000E-01			1.99	7R 2 RIBLIB	135
98413.575E-02 .064	1.000E-01		1.	1.31 2.	NB*1 RIBLIB	136
98411.806E-02 3.	2.845E 00			1.26 1.2	N9 2 RIBLIB	137
9842	2.516	2.845E 00.15			MO 2 RIBLIB	138
99401.852E-053.050E 003.050E 00				1.55	ZR 2 RIBLIB	139
99411.736E-033.050E 003.050E 00				1.36 .26	NB 2 RIBLIB	140
99422.750E 00			.92	.41 0.124	MO 2 RIBLIB	141
99432.500E-010.	0.			.1423	TC*1 RIBLIB	142
99437.665E 070.	0.	22.		.084	TC 2 RIBLIB	143
9944	0.	0.			RU 2 RIBLIB	144
100412.0A3E-016.300E 007.100E 00				3.39 1.05	NB 2 RIBLIB	145
10042	0.	0.	.2		MO 2 RIBLIB	146
100431.968E-040.	0.			1.44 .114	TC 2 RIBLIB	147
10044	0.	0.			RU 2 RIBLIB	148
101416.944E-042.500E 002.950E 00				1.03	N9 2 RIBLIB	149
101421.014E-022.500E 002.950E 00				.42 1.649	MO 2 RIBLIB	150
101439.722E-030.	0.			.47 0.362	TC 2 RIBLIB	151
10144	0.	0.			RU 2 RIBLIB	152
102427.981E-034.200E 005.990E 00				.44 1.3	MO 2 RIBLIB	153
102435.787E-050.	0.			1.79	TC 2 RIBLIB	154
10244	0.	1.4			RU 2 RIBLIB	155
103435.787E-041.550E 002.835E 00				1.03	TC 2 RIBLIB	156
103444.000E+011.450E 002.835E 00			.995	.063 0.488	RU 2 RIBLIB	157
103453.959E-020.	0.			0.040	RH*1 RIBLIB	158
10345	0.	155.	.071		RH 2 RIBLIB	159
104421.111E-031.800E 005.930E 00				1.08 1.	MO 2 RIBLIB	160
104431.250E-020.	0.			.93 3.1	TC 2 RIBLIB	161
10444	0.	0.	.7		RU 2 RIBLIB	162
104453.056E-030.	0.	800.		.0013.0002.131	RH*1 RIBLIB	163
104454.861E-040.	0.	40.		.98 0.013	RH 2 RIBLIB	164
10446	0.	0.			PD 2 RIBLIB	165
105424.630E-046.000E-012.600E 00				2.54	MO 2 RIBLIB	166
105435.556E-033.000E-011.300E 00				2.09	TC 2 RIBLIB	167
105441.846E-010.	0.	.2		.2690.42 .7680	RU 2 RIBLIB	168
105453.472E-040.	0.			0.129	RH*1 RIBLIB	169
105451.500E 000.	0.	2.100E+04.715 0.		.17 0.032	RH 2 RIBLIB	170
10546	0.	0.			PD 2 RIBLIB	171
106443.650E 023.800E-014.570E 00.1				.01	RU 2 RIBLIB	172
106450.167E-02			1.	.35 2.627	RH*1 RIBLIB	173
106453.472E-040.	0.			1.40 .222	RH 2 RIBLIB	174
10646	0.	0.			PD 2 RIBLIB	175
107431.042E-031.600E-013.600E 00				2.71	TC 2 RIBLIB	176
107442.917E-034.000E-020.				1.67 .144	RU 2 RIBLIB	177
107451.507E-020.	0.			.42 0.328	RH 2 RIBLIB	178
107462.555E 090.	0.			.014	PD 2 RIBLIB	179
10747	0.	0.	35.	0.0100.	AG 2 RIBLIB	180
108447.125E-038.000E-022.100E 00				.54 0.059	RU 2 RIBLIB	181
108451.968E-040.	0.			1.74 0.553	RH 2 RIBLIB	182
10846	0.	0.	12.2	.016	PD 2 RIBLIB	183
108471.825E 030.	0.		.9	1.605	AG*1 RIBLIB	184
108471.667E-030.	0.			.62 .017	AG 2 RIBLIB	185
10848	0.	0.	10.		CD 2 RIBLIB	186
109453.472E-042.800E-021.400E 00			1.	1.16 .31	RH 2 RIBLIB	187
109463.333E-030.	0.			0.18	PD*1 RIBLIB	188
109465.625E-010.	0.		1.	.36 .0002	PD 2 RIBLIB	189
109474.630E-040.	0.			0.088	AG*1 RIBLIB	190
10947	0.	0.	92.	.0326	AG 2 RIBLIB	191
11046	2.000E-020.	.24	.167		PD 2 RIBLIB	192
110472.600E 020.	0.	80.	0.98	.069 2.722	AG*1 RIBLIB	193
110472.778E-040.	0.			1.17 0.033	AG 2 RIBLIB	194
11048	2.000E-070.	3.1	.0322		CD 2 RIBLIB	195
111462.292E-011.000E-031.000E-02			0.32	.048 .676	PD*1 RIBLIB	196
111461.528E-021.800E-022.200E-01			1.	.85 .3	PD 2 RIBLIB	197

111478.565E-04						.065	AG*1 RIBLIB	198		
111477.500E 00					.36	0.025	AG 2 RIBLIB	199		
111483.403E-020.	0.					0.397	CO*1 RIBLIB	200		
11148	0.	0.					CO 2 RIBLIB	201		
112468.750E-011.100E-021.200E-01					.08	0.018	PD 2 RIBLIB	202		
112471.333E-010.	0.				1.44	0.695	AG 2 RIBLIB	203		
11248	0.	0.	.33	1.			CO 2 RIBLIB	204		
113461.042E-031.100E-021.000E-02					1.	1.40	PD 2 RIBLIB	205		
113478.333E-040.	0.				.1	.08	0.200	AG*1 RIBLIB	206	
113472.208E-010.	0.					.79	0.030	AG 2 RIBLIB	207	
113485.110E 030.	0.					0.999.18	.0003	CO*1 RIBLIB	208	
11348	0.	0.	2.000E 04					CO 2 RIBLIB	209	
11349			12.	.67				IN 2 RIBLIB	210	
114461.667E-031.000E-021.000E-02						.52		PD 2 RIBLIB	211	
114475.787E-050.	0.					2.02	0.5	AG 2 RIBLIB	212	
11448	0.	0.	1.24	0.1130.				CO 2 RIBLIB	213	
114495.000E 01							.192	IN*1 RIBLIB	214	
114498.333E-04						.779	.002	IN 2 RIBLIB	215	
115465.208E-04 .01	.0038				.28	1.91		PD 2 RIBLIB	216	
115472.315E-04					1.	1.49		AG*1 RIBLIB	217	
115471.458E-02 .0005	3.000E-04				.09	1.21	.2	AG 2 RIBLIB	218	
115484.300E 010.	0.				1.	.61	0.036	CO*1 RIBLIB	219	
115482.300E 000.	0.				1.	.32	0.192	CO 2 RIBLIB	220	
115491.833E-010.	0.					.055	.015	0.317	IN*1 RIBLIB	221
11549	0.	0.	203.	.78		.15		IN 2 RIBLIB	222	
11550	0.	0.						SN 2 RIBLIB	223	
116463.472E-041.000E-020.						.93		PD 2 RIBLIB	224	
116471.736E-030.	0.					2.21	1.217	AG 2 RIBLIB	225	
11648	0.	0.	1.4					CO 2 RIBLIB	226	
116493.750E-020.	0.					1.	.29	2.495	IN*1 RIBLIB	227
116491.620E-040.	0.						1.39		IN 2 RIBLIB	228
11650	3.000E-030.	.006	1.					SN 2 RIBLIB	229	
117477.639E-041.100E-021.100E-02					.5	1.63		AG 2 RIBLIB	230	
117481.208E-010.	0.				1.	.35	1.6	CO*1 RIBLIB	231	
117483.472E-020.	0.				1.	.70	0.49	CO 2 RIBLIB	232	
117497.917E-020.	0.				0.8	.52	.132	IN*1 RIBLIB	233	
117493.125E-020.	0.					.24	.73	IN 2 RIBLIB	234	
117501.400E 010.	0.						0.32	SN*1 RIBLIB	235	
11750	0.	0.						SN 2 RIBLIB	236	
118483.472E-021.000E-021.000E-03						.27	.2	CO 2 RIBLIB	237	
118495.903E-050.	0.					1.85	0.244	IN 2 RIBLIB	238	
11850	0.	0.	.31	1.				SN 2 RIBLIB	239	
119481.875E-03 .01	4.000E-03				.99	1.56		CO*1 RIBLIB	240	
119486.597E-03					1.	1.44	.1	CO 2 RIBLIB	241	
119491.250E-02					.96	1.04	0.066	IN*1 RIBLIB	242	
119491.309E-03					.1	.61	.34	IN 2 RIBLIB	243	
119502.500E+02							.089	SN*1 RIBLIB	244	
11950	0.	0.						SN 2 RIBLIB	245	
120495.787E-040.500E-020.500E-03						.88	3.2	IN 2 RIBLIB	246	
12050	0.500E-020.500E-03.141	.007						SN 2 RIBLIB	247	
121482.431E-032.500E-042.500E-03						1.73	0.85	CO 2 RIBLIB	248	
121492.153E-032.500E-042.500E-03					1.	1.58		IN*1 RIBLIB	249	
121493.472E-04 .0005	.0053					1.20	.94	IN 2 RIBLIB	250	
121509.125E 1030.500E-030.500E-02					1.	.15	.01	SN*1 RIBLIB	251	
121501.125E 001.400E-023.000E-02						.11		SN 2 RIBLIB	252	
12151	0.	0.	6.06	.01				S8 2 RIBLIB	253	
122498.681E-050.650E-020.500E-03						2.68	2.14	IN 2 RIBLIB	254	
12250	0.650E-020.500E-03.201	.005						SN 2 RIBLIB	255	
122512.847E-030.	0.						0.136	S9*1 RIBLIB	256	
122512.750E 000.	0.					.58	.405	S8 2 RIBLIB	257	
12252	0.	0.	3.	.333				TE 2 RIBLIB	258	
123491.157E-040.600E-030.500E-03						1.39	1.1	IN 2 RIBLIB	259	
123501.250E+020.600E-030.500E-03					1.	.52	0.022	SN*1 RIBLIB	260	
123502.778E-021.400E-020.						.45	0.160	SN 2 RIBLIB	261	
12351	0.	0.	3.345	.0135				S8 2 RIBLIB	262	
123521.040E+02							.249	TE*1 RIBLIB	263	

12352	0.	0.	400.					TS 2 RIBLIB	264
12450	2.000E-020.		.104	.962				SN 2 RIBLIB	265
124511.458E-020.	0.				.992	.97	0.427	S3*1 RIBLIB	266
124516.020E+010.	0.		2000.			.38	1.899	S9 2 RIBLIB	267
12452	0.	0.	7.	.714				TE 2 RIBLIB	268
125506.736E-031.100E-021.000E-03					1.	.79	0.361	SN*1 RIBLIB	269
125509.400E 001.200E-027.100E-02						.92	0.094	SN 2 RIBLIB	270
125519.855E 020.	0.		20.		.26	.096	.415	S9 2 RIBLIB	271
125525.800E 010.	0.						0.145	TE*1 RIBLIB	272
12552	0.	0.	1.5					TE 2 RIBLIB	273
126503.650E+071.000E-012.000E-01					1.	.12	.09	SN 2 RIBLIB	274
126511.319E-020.	0.				.01	.00780	.117	SE*1 RIBLIB	275
126511.250E 010.	0.					.74	1.780	S9 2 RIBLIB	276
12652	0.	0.	1.	.1				TE 2 RIBLIB	277
127509.750E-022.400E-013.900E-01						1.34	1.	SN 2 RIBLIB	278
127513.900E 001.000E-021.000E-02					.2	.38	0.505	S9 2 RIBLIB	279
127521.050E 020.	0.				.015	.00410	.088	TE*1 RIBLIB	280
127523.875E-010.	0.					.23	0.004	TE 2 RIBLIB	281
12753	0.	0.	6.4					I 2 RIBLIB	282
128504.306E-023.700E-014.000E-01					.03	3.85	3.27	SN 2 RIBLIB	283
128514.000E-015.000E-020.					1.	.35	3.6	S8*1 RIBLIB	284
128516.944E-039.000E-021.000E-01						1.20	1.7	S8 2 RIBLIB	285
12852	0.	0.	.157	.108				TE 2 RIBLIB	286
128531.736E-023.000E-052.000E-04						.77	0.093	I 2 RIBLIB	287
12854	0.	0.	5.					XE 2 RIBLIB	288
129511.792E-011.000E 001.000E 00					.36	.542	.916	S3 2 RIBLIB	289
129523.300E 010.	0.						0.106	TE*1 RIBLIB	290
129524.653E-020.	0.					.45	.238	TE 2 RIBLIB	291
129535.840E 090.	0.		28.			.050	.040	I 2 RIBLIB	292
12954	0.	0.	25.					XE 2 RIBLIB	293
130501.800E-032.000E 002.000E 00						4.23		SN 2 RIBLIB	294
130512.292E-020.	0.					1.72	1.0	S3 2 RIBLIB	295
13052	0.	0.	.23	.13				TE 2 RIBLIB	296
130535.200E-01 .0005 .0052 18.						.27	2.123	I 2 RIBLIB	297
13054	0.	0.	5.					XE 2 RIBLIB	298
131511.597E-022.700E 003.100E 00					.15	1.33		S9 2 RIBLIB	299
131521.200E 004.000E-021.300E-01					.783	.16	1.468	TE*1 RIBLIB	300
131521.736E-021.600E-015.400E-01						.74	0.376	TE 2 RIBLIB	301
131539.050E 00 .2	0.		.7		.007	.18	0.399	I 2 RIBLIB	302
131541.200E 010.	1.000E-02						0.164	XE*1 RIBLIB	303
13154	0.	0.	85.					XE 2 RIBLIB	304
132511.458E-033.400E 005.100E 00						2.68		S8 2 RIBLIB	305
132523.250E 001.000E 00 .18						.055	0.280	TE 2 RIBLIB	306
132539.583E-020.	0.					.49	2.249	I 2 RIBLIB	307
13254	0.	0.	5.2	.962				XE 2 RIBLIB	308
133512.847E-033.800E 003.000E 00					.72	1.72		S8 2 RIBLIB	309
133523.472E-021.100E 001.000E 00					.87	.49	1.613	TE*1 RIBLIB	310
133521.389E-031.100E 001.000E 00						.97	0.6	TE 2 RIBLIB	311
133538.750E-015.000E-012.000E-01					.02	.46	.53	I 2 RIBLIB	312
133542.300E 000.	0.						0.233	XE*1 RIBLIB	313
133545.270E 000.	1.700E 00190.					.10	.082	XE 2 RIBLIB	314
13355	0.	0.	30.6	.085				CS 2 RIBLIB	315
134515.556E-043.000E 003.500E 00						3.45	0.290	S3 2 RIBLIB	316
134522.917E-023.700E 004.000E 00						.52	.1	TE 2 RIBLIB	317
134533.681E-029.000E-010.						.73	2.5	I 2 RIBLIB	318
13454	0.	0.	5.2	.962				XE 2 RIBLIB	319
134551.200E-010.	0.				.01	.00170	.153	CS*1 RIBLIB	320
134557.665E 020.	0.		136.			.16	1.588	CS 2 RIBLIB	321
13456	0.	0.	4.16	.038				BA 2 RIBLIB	322
135512.779E-04 4.3	1.1					3.16		S8 2 RIBLIB	323
135526.944E-04						2.68		TE 2 RIBLIB	324
135532.792E-01 1.9	5.700E 00				.27	.32	1.642	I 2 RIBLIB	325
135541.111E-020.	0.						0.530	XE*1 RIBLIB	326
135543.833E-013.000E-01 .47	2.900E+06					.31	0.261	XE 2 RIBLIB	327
135557.300E+080.			8.7			.055		CS 2 RIBLIB	328
135561.200E 000.	0.						0.268	BA*1 RIBLIB	329

13556	0.	0.	5.				BA 2	RIBLIB	330
136539.606E-043.100E 002.100E 00				1.68	3.237		I 2	RIBLIB	331
13654	3.76	4.520E 00.15					XE 2	RIBLIB	332
136551.300E 016.900E-031.100E-01				.11	2.466		CS 2	RIBLIB	333
13656	0.	0.	1.01	.01			BA 2	RIBLIB	334
137532.774E-044.900E 005.100E 00				2.19	.71		I 2	RIBLIB	335
137542.708E-031.300E 001.530E 00				1.48	1.		XE 2	RIBLIB	336
137551.095E+040.	0.		.11	.92	.20		CS 2	RIBLIB	337
137561.806E-030.	0.					0.662	BA*1	RIBLIB	338
13756	0.	0.	4.				BA 2	RIBLIB	339
138537.292E-053.400E 003.700E 00				3.10	1.5		I 2	RIBLIB	340
138549.722E-032.100E 002.210E 00				.96	0.420		XE 2	RIBLIB	341
138552.236E-023.000E-014.000E-01				1.10	2.153		CS 2	RIBLIB	342
13856	0.	0.	.4				BA 2	RIBLIB	343
139532.314E-051.800E 001.700E 00				2.40			I 2	RIBLIB	344
139544.745E-042.900E 002.800E 00				2.00	.4		XE 2	RIBLIB	345
139556.597E-031.200E 001.230E 00				1.74	0.232		CS 2	RIBLIB	346
139565.764E-021.000E-011.400E-014.				.91	.053		BA 2	RIBLIB	347
13957	0.	0.	8.9				LA 2	RIBLIB	348
140541.852E-043.700E 003.300E 00				1.09			XE 2	RIBLIB	349
140557.639E-042.300E 002.100E 00				2.66	.59		CS 2	RIBLIB	350
140561.280E 013.000E-010.			20.	.30	.222		BA 2	RIBLIB	351
140571.675E 000.		2.000E-01		.48	2.468		LA 2	RIBLIB	352
14058	0.	0.	.6				CE 2	RIBLIB	353
141541.068E-051.800E 001.100E 00				2.36			XE 2	RIBLIB	354
141552.773E-042.900E 003.700E 00				1.72			CS 2	RIBLIB	355
141561.250E-021.200E 009.000E-01				1.14	.46		BA 2	RIBLIB	356
141571.625E-011.000E-010.				.96	.0272		LA 2	RIBLIB	357
141583.250E 010.	0.		30.	.15	0.101		CE 2	RIBLIB	358
14159	0.	0.	12.				PR 2	RIBLIB	359
142541.736E-053.500E-013.100E-01				1.70			XE 2	RIBLIB	360
142552.662E-053.050E 002.870E 00				2.98			CS 2	RIBLIB	361
142567.639E-032.200E 001.830E 00				.78	.112		BA 2	RIBLIB	362
142575.833E-023.000E-010.				1.42	1.164		LA 2	RIBLIB	363
14258	0.	0.	1.				CE 2	RIBLIB	364
142598.000E-010.	0.		20.	.83	.063		PR 2	RIBLIB	365
14260	0.	0.	17.				NO 2	RIBLIB	366
143541.157E-053.100E 002.650E 00				2.91			XE 2	RIBLIB	367
143552.314E-053.100E 002.650E 00				2.32			CS 2	RIBLIB	368
143561.789E-04				1.47			BA 2	RIBLIB	369
143579.722E-03				1.33	1.		LA 2	RIBLIB	370
143581.375E 000.	0.		6.	.37	.377		CE 2	RIBLIB	371
143591.370E 010.	0.		89.	.32			PR 2	RIBLIB	372
14360	0.	0.	330.				NO 2	RIBLIB	373
144541.157E-052.900E 001.600E 00				2.19			XE 2	RIBLIB	374
144552.315E-052.900E 001.600E 00				3.50			CS 2	RIBLIB	375
144563.472E-05				1.00			BA 2	RIBLIB	376
144574.630E-05				2.23			LA 2	RIBLIB	377
144582.850E 023.000E-015.900E-011.				.080	0.030		CE 2	RIBLIB	378
144591.201E-020.		1.400E-01		1.22	.031		PR 2	RIBLIB	379
14460	0.	0.	5.				NO 2	RIBLIB	380
145582.083E-034.200E 003.130E 00				.77			CE 2	RIBLIB	381
145592.458E-010.	0.			.68	.056		PR 2	RIBLIB	382
14560	0.	0.	50.				NO 2	RIBLIB	383
146589.722E-033.200E 002.600E 00				.22	0.320		CE 2	RIBLIB	384
146591.667E-021.000E-010.				1.27	1.107		PR 2	RIBLIB	385
14660	0.	0.	2.				NO 2	RIBLIB	386
147588.333E-040.650E 000.550E 00				1.55			CE 2	RIBLIB	387
147598.333E-030.650E 000.550E 00				1.12	1.		PR 2	RIBLIB	388
147601.110E+011.300E 001.100E 00				.23	0.196		NO 2	RIBLIB	389
147619.855E 020.	0.		230.	.522	.059		PM 2	RIBLIB	390
14762	0.	0.	90.				SM 2	RIBLIB	391
148584.861E-041.710E 001.730E 00				.95			CE 2	RIBLIB	392
148591.354E-030.	0.			2.17	0.300		PR 2	RIBLIB	393
14860	0.	0.	4.				NO 2	RIBLIB	394
148615.400E 000.	0.		2000.	1.	.39	1.343	PM*1	RIBLIB	395

148614.100E 010.	0.	3.000E+04	.41	1.227	PM 2 RIBLIB	396
14862	0.	0.			SM 2 RIBLIB	397
149607.500E-021.300E 001.320E 00			.55	.5	ND 2 RIBLIB	398
149612.208E 000.	0.		.36	.J35	PM 2 RIBLIB	399
14962	0.	4.150E 04			SM 2 RIBLIB	400
15060	7.400E-011.010E 001.5				ND 2 RIBLIB	401
15062	0.	100.			SM 2 RIBLIB	402
151608.333E-034.800E-018.000E-01			.63	0.717	ND 2 RIBLIB	403
151611.183E 002.000E-020.			.38	.715	PM 2 RIBLIB	404
151623.285E 040.	0.	1.500E 04	.019	.0004	SM 2 RIBLIB	405
15163	0.	3.700E 03.322			EU 2 RIBLIB	406
152614.167E-031.500E-013.100E-01			.86	2.8	PM 2 RIBLIB	407
15262	1.500E-013.100E-01210.				SM 2 RIBLIB	408
152633.875E-010.	0.		.60	.238	EU*1 RIBLIB	409
152634.745E 030.	0.	5000.	.19	1.209	EU 2 RIBLIB	410
15264	0.	180.			GD 2 RIBLIB	411
153613.819E-03 .075	.185.		.62	.3	PM 2 RIBLIB	412
153621.958E 00 .075	.185		.22	.108	SM 2 RIBLIB	413
15363	0.	320.			EU 2 RIBLIB	414
154611.736E-03 .045	.145		1.		PM 2 RIBLIB	415
15462	.045	.145	5.		SM 2 RIBLIB	416
154635.840E 030.	0.	1400.	.20	1.351	EU 2 RIBLIB	417
15464	0.	0.			GD 2 RIBLIB	418
155621.528E-023.100E-022.300E-01			.60	0.115	SM 2 RIBLIB	419
155636.570E 020.	0.	1.300E 04	.046	0.072	EU 2 RIBLIB	420
15564	0.	5.800E 04			GD 2 RIBLIB	421
156623.917E-011.300E-021.100E-01			.30		SM 2 RIBLIB	422
156631.500E 010.	0.		.70	1.2	EU 2 RIBLIB	423
15664	0.	0.			GD 2 RIBLIB	424
157636.417E-017.400E-037.400E-02			.42	0.525	EU 2 RIBLIB	425
15764	0.	2.400E 05			GD 2 RIBLIB	426
158633.104E-022.000E-034.000E-02			1.04	0.600	EU 2 RIBLIB	427
15864	0.	3.4			GD 2 RIBLIB	428
159631.319E-02.00055	.0105		.86	.2	EU 2 RIBLIB	429
159647.500E-01.00055	.0105		.29	0.075	GD 2 RIBLIB	430
15965	0.	46.			TA 2 RIBLIB	431
160631.736E-02.00135	.00081		1.5		EU 2 RIBLIB	432
16064	.00135	.00081	.8		GD 2 RIBLIB	433
160657.300E 010.	0.	525.	.20	1.193	TA 2 RIBLIB	434
16066	0.	0.			OY 2 RIBLIB	435
161642.569E-03 .008	3.900E-03		.59	0.428	GD 2 RIBLIB	436
161656.900E 000.	0.		.15	0.037	TB 2 RIBLIB	437
16166	0.	580.			OY 2 RIBLIB	438
162643.650E+02	1.800E-03		.34	1.	GD 2 RIBLIB	439
162659.333E-02			1.68		T9 2 RIBLIB	440
16266		140.			OY 2 RIBLIB	441
163652.917E-01	4.000E-04		1.22	.2	T8 2 RIBLIB	442
16366	4.000E-04120.				OY 2 RIBLIB	443
16466	3.000E-042800.	.714			OY 2 RIBLIB	444
165669.727E-040.	1.200E-04		.03	.00840.120	OY*1 RIBLIB	445
165669.583E-020.	0.	4700.	.44	.0269	OY 2 RIBLIB	446
16567	0.	65.			HO 2 RIBLIB	447
166663.333E 000.	6.800E-05		.12	0.088	OY 2 RIBLIB	448
166671.133E 000.	0.		.67	.0558	HO 2 RIBLIB	449
16668	0.	12.	.01		ER 2 RIBLIB	450
167692.892E-050.	0.			0.208	ER*1 RIBLIB	451
16769		700.			ER 2 RIBLIB	452
145					RIBLIB	453
3 1 4500.					H 2 RIBLIB	454
7 4 53.6					BE 2 RIBLIB	455
14 6 2.0 E6					C 2 RIBLIB	456
18 9 0.078					F 2 RIBLIB	457
2211 950.					NA 2 RIBLIB	458
2411 0.63					NA 2 RIBLIB	459
3114 0.11					SI 2 RIBLIB	460
3215 14.3					P 2 RIBLIB	461

3516	87.1	
3617	1.2	E8
3817	0.025	
4118	0.0763	
4219	.52	
4520	16.	
4720	4.	
4621	85.	
4721	3.47	
4821	1.83	
4823	16.1	
5124	27.8	
5225	5.55	
5425	300.	
5625	.11	
5526	1100.	
5926	45.1	
5727	270.	
5827	0.38	
5827	72.	
6027	1.9	E3
5928	2.9	E7
6328	2.9	E4
6528	.11	
6429	.53	
6530	245.	
6930	.58	
6930	.036	
7132	12.	
7333	76.	
7433	17.5	
7534	127.	
8538	.049	
8538	65.	
9643	.036	
9643	4.3	
9743	.25	
9743	3.7	E6
9744	2.8	
10346	17.	
10547	40.	
10948	475.	
11349	.073	
11350	112.	
12653	13.3	
13155	10.	
13156	11.6	
15364	236.	
16968	9.4	
17168	0.31	
17069	127.	
17169	694.	
17570	4.1	
17771	6.8	
18172	46.	
18273	112.	
18174	140.	
18574	74.	
18774	1.	
18375	73.	
18675	3.79	
18775	1.8E13	
18875	.71	
18576	95.	
19176	.58	
19176	16.	

S	2	RIBLIB	462
CL	2	RIBLIB	463
CL	2	RIBLIB	464
AR	2	RIBLIB	465
K	2	RIBLIB	466
CA	2	RIBLIB	467
CA	2	RIBLIB	468
SC	2	RIBLIB	469
SC	2	RIBLIB	470
SC	2	RIBLIB	471
V	2	RIBLIB	472
CR	2	RIBLIB	473
MN	2	RIBLIB	474
MN	2	RIBLIB	475
MN	2	RIBLIB	476
FE	2	RIBLIB	477
FE	2	RIBLIB	478
CO	2	RIBLIB	479
CO*	1	RIBLIB	480
CO	2	RIBLIB	481
CO	2	RIBLIB	482
NI	2	RIBLIB	483
NI	2	RIBLIB	484
NI	2	RIBLIB	485
CU	2	RIBLIB	486
ZN	2	RIBLIB	487
ZN*	1	RIBLIB	488
ZN	2	RIBLIB	489
GE	2	RIBLIB	490
AS	2	RIBLIB	491
AS	2	RIBLIB	492
SE	2	RIBLIB	493
SR*	1	RIBLIB	494
SR	2	RIBLIB	495
TC*	1	RIBLIB	496
TC	2	RIBLIB	497
TC*	1	RIBLIB	498
TC	2	RIBLIB	499
RU	2	RIBLIB	500
PD	2	RIBLIB	501
AG	2	RIBLIB	502
CD	2	RIBLIB	503
IN*	1	RIBLIB	504
SN	2	RIBLIB	505
I	2	RIBLIB	506
CS	2	RIBLIB	507
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GD	2	RIBLIB	509
ER	2	RIBLIB	510
ER	2	RIBLIB	511
TM	2	RIBLIB	512
TM	2	RIBLIB	513
YB	2	RIBLIB	514
IU	2	RIBLIB	515
HF	2	RIBLIB	516
TA	2	RIBLIB	517
W	2	RIBLIB	518
W	2	RIBLIB	519
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RE	2	RIBLIB	521
RE	2	RIBLIB	522
RE	2	RIBLIB	523
RE	2	RIBLIB	524
OS	2	RIBLIB	525
OS*	1	RIBLIB	526
OS	2	RIBLIB	527

19376	1.3
19077	12.
19277	74.5
19477	.79
19178	3.
19378	3.4
19378	1.8 E5
19778	.058
19779	.75
19679	5.6
19879	2.7
19979	3.15
19780	1.
19780	2.7
20380	45.3
20081	1.13
20181	3.
20281	12.
20481	1.1 E3
20382	2.17
21082	7.1 F3
21282	.44
20683	6.4
20783	2.9 F3
21083	5.0
21283	.042
21084	138.4
21185	.3
22388	11.7
22488	3.64
22688	5.9 F5
22888	2.4 F3
22789	8.0 E3
22889	.26
22790	18.4
22890	700.
23090	2.9 E7
23190	1.07
23290	5.1E12
23490	24.1
23091	17.7
23191	1.3 E7
23391	27.4
23092	20.8
23292	2.7 F4
23392	5.9 E7
23492	9.1 E7
23592	2.6E11
23692	8.7 E9
23892	1.7E12
23793	8.0 E8
23993	2.33
23894	3.3 E4
23994	8.9 E6
24094	2.4 E6
24194	4.8 E3
24294	1.4 E8
24195	1.7 E5
24295	5.6 E4
24295	0.677
24395	2.9 E6
24296	162.5
24396	1.3 E4
24496	6.7 E3
24596	7.3 E6
24696	2.4 E6

OS 2	RIBLIB	529
IR 2	RIBLIB	529
IR 2	RIBLIB	530
IR 2	RIBLIB	531
PT 2	RIBLIB	532
PT*1	RIBLIB	533
PT 2	RIBLIB	534
PT*1	RIBLIB	535
PT 2	RIBLIB	536
AU 2	RIBLIB	537
AU 2	RIBLIB	539
AU 2	RIBLIB	539
HG*1	RIBLIB	540
HG 2	RIBLIB	541
HG 2	RIBLIB	542
TL 2	RIBLIB	543
TL 2	RIBLIB	544
TL 2	RIBLIB	545
TL 2	RIBLIB	546
PB 2	RIBLIB	547
PB 2	RIBLIB	548
PB 2	RIBLIB	549
BI 2	RIBLIB	550
BI 2	RIBLIB	551
BI 2	RIBLIB	552
BI 2	RIBLIB	553
PO 2	RIBLIB	554
AT 2	RIBLIB	555
RA 2	RIBLIB	556
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RA 2	RIBLIB	559
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PU 2	RIBLIB	581
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PU 2	RIBLIB	583
PU 2	RIBLIB	584
AM 2	RIBLIB	585
AM*1	RIBLIB	586
AM 2	RIBLIB	587
AM 2	RIBLIB	588
CM 2	RIBLIB	589
CM 2	RIBLIB	590
CM 2	RIBLIB	591
CM 2	RIBLIB	592
CM 2	RIBLIB	593

24997 290.
24998 1.31 E5
25098 3.7 E3
25298 803.

BK 2 RIBLIB 594
CF 2 RIALIB 595
CF 2 RIBLIB 596
CF 2 RIALIB 597
RIALIB 598

ORGAN DATA LIBRARY

1136 ORGAN DATA LIBRARY VERSION 1A SEPTEMBER 4, 1974									
1	3	H	1	.450+04	.120+02	.100+01	.100+01	.100+01	.100-01
1	3	H	2	.450+04	.120+02	.100+01	.100+01	.100+01	.100-01
1	3	H	8						.100-01
4	7	EE	1	.536+02	.180+03	.200-02	.250-00	.100+01	.350-01
4	7	EE	3	.536+02	.120+03	.600-04	.750-02	.300-01	.120-01
4	7	EE	4	.536+02	.270+03	.200-03	.250-01	.100+00	.160-01
4	7	EE	5	.536+02	.540+03	.400-05	.500-03	.200-02	.120-01
4	7	EE	6	.536+02	.450+03	.640-03	.800-01	.320-00	.350-01
4	7	EE	8						.160-01
6	14	C	1	.200+07	.100+02	.100+01	.750-00	.100+01	.540-01
6	14	C	6	.200+07	.400+02	.250-01	.200-01	.250-01	.270-00
6	14	C	7	.200+07	.120+02	.500-00	.380-00	.500-00	.540-01
9	18	F	1	.780-01	.858+03	.100+01	.750-00	.100+01	.890-00
9	18	F	6	.780-01	.145+04	.530-00	.400-00	.530-00	.140+01
9	18	F	8						.540+00
11	22	NA	1	.350+03	.110+02	.100+01	.750-00	.100+01	.160+01
11	22	NA	8						.300+00
11	24	NA	1	.630-00	.110+02	.100+01	.750-00	.100+01	.270+01
11	24	NA	6						.150+01
14	31	SI	1	.110+00	.600+02	.650-00	.660-00	.100+01	.590-00
14	31	SI	8	.110+00	.600+02	.900-01	.700-01	.100+00	.590-00
14	31	SI	9	.110+00	.600+02	.850-03	.700-03	.100-02	.530-00
14	31	SI10		.110+00	.600+02	.430-03	.340-03	.500-03	.590-00
14	31	SI11		.110+00	.600+02	.500-04	.400-04	.600-03	.590-00
14	31	SI12		.110+00	.600+02	.300-01	.300-01	.400-01	.590-00
15	32	P	1	.143+02	.257+03	.750-00	.630-00	.100+01	.690-00
15	32	P	4	.143+02	.120+02	.500-01	.400-01	.700-01	.690-00
15	32	P	6	.143+02	.115+04	.375-00	.320-00	.200-00	.350+01
15	32	P	8						.690+00
15	32	P13		.143+02	.257+03	.530-02	.440-02	.700-02	.690-00
16	35	S	1	.371+02	.900+02	.100+01	.750-00	.100+01	.560-01
16	35	S	6	.371+02	.600+03	.300-01	.200-01	.300-01	.280-00
16	35	S	8						.560-01
16	35	S10		.371+02	.623+03	.130-02	.960-03	.130-02	.560-01
16	35	S12		.371+02	.153+04	.100-01	.750-02	.100-01	.560-01
17	36	OL	1	.120+09	.290+02	.100+01	.750-00	.100+01	.260-00
17	36	OL	8						.260+00
17	36	OL	1	.260-01	.290+02	.100+01	.750-00	.100+01	.230+01
17	36	OL	8						.160+01
19	42	K	1	.520-00	.580+02	.100+01	.750-00	.100+01	.160+01
19	42	K	4	.520-00	.580+02	.200-01	.150-01	.200-01	.150+01
19	42	K	5	.520-00	.580+02	.400-02	.300-02	.400-02	.150+01
19	42	K	8						.150+01
19	42	K13		.520-00	.580+02	.400-01	.300-01	.400-01	.150+01
19	42	K14		.520-00	.580+02	.050-00	.490-00	.650-00	.160+01
20	45	CA	1	.164+03	.164+03	.500-00	.550-00	.100+01	.360-01
20	45	CA	6	.164+03	.164+03	.540-00	.500-00	.900-00	.430-00
20	45	CA	8						.860-01
20	47	CA	1	.490+01	.164+03	.600-00	.550-00	.100+01	.140+01
20	47	CA	6	.490+01	.160+03	.540-00	.500-00	.900-00	.260+01
20	47	CA	8						.657-00.81-00.52-00
21	46	SC	1	.850+02	.300+02	.100-03	.250-00	.100+01	.130+01
21	46	SC	3	.850+02	.750+02	.200-05	.500-02	.200-01	.500-00
21	46	SC	4	.850+02	.300+02	.150-04	.400-01	.150-00	.640-00
21	46	SC	6	.850+02	.330+02	.200-04	.500-01	.200-00	.900-00
21	46	SC	8						.640+00
21	47	SC	1	.343+01	.300+02	.100-03	.250-00	.100+01	.260-00
21	47	SC	3	.343+01	.750+02	.200-05	.500-02	.200-01	.200-00
21	47	SC	4	.343+01	.300+02	.150-04	.400-01	.150-00	.210-00
21	47	SC	6	.343+01	.330+02	.200-04	.500-01	.200-00	.290-00
21	47	SC	8						.210+00
21	48	SC	1	.183+01	.300+02	.100-03	.250-00	.100+01	.220+01

21 46 SC 4	.183+01	.369+02	.150-04	.400-01	.150-00	.110+01
21 48 SC 6	.183+01	.330+02	.200-04	.500-01	.200-00	.160+01
21 48 SC 8						.110+01
23 48 V 1	.161+02	.420+02	.200-01	.260-00	.190+01	.190+01
23 48 V 3	.161+02	.740+02	.600-03	.100-01	.400-01	.700-00
23 48 V 4	.161+02	.700+02	.120-02	.200-01	.600-01	.900-00
23 48 V 5	.161+02	.900+02	.200-03	.260-02	.100-01	.700-00
23 48 V 6	.161+02	.135+03	.280-02	.400-01	.140-00	.120+01
23 48 V 8						.900+00
24 51 CR 1	.278+02	.616+03	.500-02	.250-00	.100+01	.250-01
24 51 CR 3	.278+02	.616+03	.130-04	.660-03	.270-02	.120-01
24 51 CR 6	.278+02	.616+03	.200-03	.100-01	.400-01	.140-01
24 51 CR 15	.278+02	.616+03	.450-05	.230-03	.900-03	.340-02
24 51 CR 16	.278+02	.616+03	.450-05	.230-03	.900-03	.840-02
25 52 MN 1	.555+01	.170+02	.100+00	.300-00	.100+01	.210+01
25 52 MN 4	.555+01	.250+02	.200-01	.700-01	.240-00	.960-00
25 52 MN 8						.960+00
25 52 MN 17	.555+01	.570+01	.300-02	.900-02	.300-01	.560-00
25 54 MN 1	.300+03	.170+02	.100+00	.300-00	.100+01	.510-00
25 54 MN 4	.300+03	.250+02	.200-01	.700-01	.240-00	.230-00
25 54 MN 8						.230+00
25 54 MN 17	.300+03	.570+01	.300-02	.900-02	.300-01	.130-00
25 56 MN 1	.110+00	.170+02	.100+00	.300-00	.100+01	.190+01
25 56 MN 4	.110+00	.250+02	.200-01	.700-01	.240-00	.130+01
25 56 MN 8						.130+01
25 56 MN 17	.110+00	.570+01	.300-02	.900-02	.300-01	.110+01
26 55 FE 1	.110+04	.800+03	.100+00	.300-00	.100+01	.650-02
26 55 FE 4	.110+04	.550+03	.130-01	.400-01	.130-00	.650-02
26 55 FE 5	.110+04	.600+03	.200-02	.600-02	.200-01	.650-02
26 55 FE 6	.110+04	.160+04	.100-01	.300-01	.100+00	.650-02
26 55 FE 8	.110+04	.320+04	.200-02	.600-02	.200-01	.650-02
26 59 FE 1	.451+02	.800+03	.100+00	.300-00	.100+01	.310-00
26 59 FE 4	.451+02	.550+03	.130-01	.400-01	.130-00	.420-00
26 59 FE 5	.451+02	.600+03	.200-02	.600-02	.200-01	.340-00
26 59 FE 8	.451+02	.320+04	.200-02	.600-02	.200-01	.420-00
27 57 CO 1	.270+03	.950+01	.300-00	.400-00	.100+01	.900-01
27 57 CO 4	.270+03	.950+01	.700-02	.200-01	.400-01	.530-01
27 57 CO 5	.270+03	.950+01	.420-03	.560-03	.140-02	.450-01
27 57 CO 8						.530-01
27 57 CO 17	.270+03	.950+01	.600-03	.800-03	.200-02	.400-01
27 58 CO 1	.720+02	.950+01	.300-00	.400-00	.100+01	.610-00
27 58 CO 4	.720+02	.950+01	.700-02	.200-01	.400-01	.290-00
27 58 CO 5	.720+02	.950+01	.420-03	.560-03	.140-02	.220-00
27 58 CO 6						.290+00
27 58 CO 17	.720+02	.950+01	.600-03	.800-03	.200-02	.170-00
27 58100 1	.380-00	.950+01	.300-00	.400-00	.100+01	.900-01
27 58100 4	.380-00	.950+01	.700-02	.200-01	.400-01	.590-01
27 58100 5	.380-00	.950+01	.420-03	.560-03	.140-02	.480-01
27 58100 8						.230-01
27 58100 17	.380-00	.950+01	.600-03	.800-03	.200-02	.390-01
27 60 CO 1	.190+04	.950+01	.300-00	.400-00	.100+01	.150+01
27 60 CO 4	.190+04	.950+01	.700-02	.200-01	.400-01	.720-00
27 60 CO 5	.190+04	.950+01	.420-03	.560-03	.140-02	.560-00
27 60 CO 8						.720+00
27 60 CO 17	.190+04	.950+01	.600-03	.800-03	.200-02	.440-00
28 59 NI 1	.290+08	.667+03	.300-00	.400-00	.100+01	.770-02
28 59 NI 4	.290+08	.500+03	.200-01	.300-01	.700-01	.770-02
28 59 NI 6	.290+08	.800+03	.150-00	.200-00	.500-00	.770-02
28 59 NI 8						.770-02
28 63 NI 1	.290+05	.667+03	.300-00	.400-00	.100+01	.210-01
28 63 NI 4	.290+05	.500+03	.200-01	.300-01	.700-01	.210-01
28 63 NI 6	.290+05	.300+03	.150-00	.200-00	.500-00	.110+00
28 63 NI 8						.210-01
28 65 NI 1	.110+00	.667+03	.300-00	.400-00	.100+01	.140+01
28 65 NI 4	.110+00	.500+03	.200-01	.300-01	.700-01	.120+01

28 65 NI 6							.120+01
29 64 CU 1	.530-00	.900+02	.280-00	.390-00	.100+01	.250-00	
29 64 CU 3	.530-00	.100+02	.100-01	.200-01	.500-01	.170-00	
29 64 CU 4	.530-00	.150+03	.200-01	.300-01	.800-01	.190-00	
29 64 CU 5	.530-00	.200+01	.200-01	.300-01	.700-01	.170-00	
29 64 CU 6						.190+00	
29 64 CU13	.530-00	.800+03	.300-02	.400-02	.100-01	.210-00	
29 64 CU18	.530-00	.800+02	.300-02	.400-02	.100-01	.170-00	
30 65 ZN 1	.245+03	.933+03	.100+00	.300-00	.100+01	.320-00	
30 65 ZN 3	.245+03	.149+03	.400-02	.120-01	.400-01	.110+00	
30 65 ZN 4	.245+03	.910+02	.350-01	.110+00	.350-00	.150-00	
30 65 ZN 6	.245+03	.130+04	.150-01	.450-01	.150-00	.940-01	
30 65 ZN 8						.150+00	
30 65 ZN10	.245+03	.270+03	.900-04	.270-03	.900-03	.560-01	
30 65 ZN11	.245+03	.107+03	.400-04	.120-03	.400-03	.560-01	
30 65 ZN14	.245+03	.196+04	.300-01	.900-01	.300-00	.320-00	
30 65 ZN15	.245+03	.140+02	.600-02	.200-01	.600-01	.560-01	
30 65 ZN17	.245+03	.250+02	.300-02	.900-02	.300-01	.640-01	
30 69 ZN 1	.360-01	.933+03	.100+00	.300-00	.100+01	.370-00	
30 69 ZN 3	.360-01	.149+03	.400-02	.120-01	.400-01	.370-00	
30 69 ZN 4	.360-01	.910+02	.350-01	.110+00	.350-00	.370-00	
30 69 ZN 6	.360-01	.130+04	.150-01	.450-01	.150-00	.190+01	
30 69 ZN 8						.370+00	
30 69 ZN10	.360-01	.270+03	.900-04	.270-03	.900-03	.370-00	
30 69 ZN11	.360-01	.107+03	.400-04	.120-03	.400-03	.370-00	
30 69 ZN14	.360-01	.196+04	.300-01	.900-01	.300-00	.370-00	
30 69 ZN15	.360-01	.140+02	.600-02	.200-01	.600-01	.370-00	
30 69 ZN17	.360-01	.250+02	.300-02	.900-02	.300-01	.370-00	
30 691ZN 1	.580-00	.933+03	.100+00	.300-00	.100+01	.640-00	
30 691ZN 3	.580-00	.149+03	.400-02	.120-01	.400-01	.470-00	
30 691ZN 4	.580-00	.910+02	.350-01	.110+00	.350-00	.500-00	
30 691ZN 6	.580-00	.130+04	.150-01	.450-01	.150-00	.210+01	
30 691ZN 8						.467-00	.50-00 .50-00
30 691ZN10	.580-00	.270+03	.900-04	.270-03	.900-03	.430-00	
30 691ZN11	.580-00	.107+03	.400-04	.120-03	.400-03	.430-00	
30 691ZN14	.580-00	.196+04	.300-01	.900-01	.300-00	.640-00	
30 691ZN15	.580-00	.140+02	.600-02	.200-01	.600-01	.430-00	
30 691ZN17	.580-00	.250+02	.300-02	.900-02	.300-01	.450-00	
31 72 GA 1	.590-00	.600+01	.100-02	.250-00	.100+01	.180+01	
31 72 GA 3	.590-00	.900+01	.200-04	.500-02	.200-01	.890-00	
31 72 GA 4	.590-00	.480+01	.250-03	.630-01	.250-00	.110+01	
31 72 GA 5	.590-00	.600+01	.100-04	.250-02	.100-01	.390-00	
31 72 GA 6	.590-00	.120+02	.300-03	.750-01	.300-00	.260+01	
31 72 GA 8						.110+01	
32 71 GE 1	.120+02	.100+01	.100-01	.260-00	.100+01	.100-01	
32 71 GE 3	.120+02	.120+02	.300-03	.800-02	.300-01	.100-01	
32 71 GE 4	.120+02	.750+01	.200-03	.500-02	.200-01	.100-01	
32 71 GE 8						.190-01	
33 73 AS 1	.760+02	.260+03	.300-01	.270-00	.100+01	.610-01	
33 73 AS 3	.760+02	.550+03	.300-03	.270-02	.100-01	.360-01	
33 73 AS 4	.760+02	.550+03	.900-03	.800-02	.300-01	.410-01	
33 73 AS 8						.410-01	
33 74 AS 1	.175+02	.200+03	.300-01	.270-00	.100+01	.560-00	
33 74 AS 3	.175+02	.550+03	.300-03	.270-02	.100-01	.340-00	
33 74 AS 4	.175+02	.550+03	.900-03	.800-02	.300-01	.300-00	
33 74 AS 8						.380+00	
33 76 AS 1	.111+01	.290+03	.300-01	.270-00	.100+01	.130+01	
33 76 AS 3	.111+01	.550+03	.300-03	.270-02	.100-01	.110+01	
33 76 AS 4	.111+01	.550+03	.900-03	.800-02	.300-01	.110+01	
33 76 AS 8						.110+01	
33 77 AS 1	.162+01	.260+03	.300-01	.270-00	.100+01	.240-00	
33 77 AS 3	.162+01	.550+03	.300-03	.270-02	.100-01	.240-00	
33 77 AS 4	.162+01	.550+03	.900-03	.800-02	.300-01	.240-00	
33 77 AS 8						.240+00	
34 75 SE 1	.127+03	.110+02	.900-00	.700-00	.100+01	.200-00	

34 75	SR 4	.127+03	.240+02	.600-01	.500-01	.700-01	.940-01
34 75	SR 5	.127+03	.180+02	.450-02	.350-02	.500-02	.720-01
34 75	SR 8						.940+01
35 82	SR 1	.150+01	.860+01	.100+01	.750-00	.100+01	.180+01
35 82	SR 8						.850+00
37 86	SR 1	.186+02	.450+02	.100+01	.750-00	.150+01	.700-00
37 86	SR 4	.186+02	.630+02	.500-01	.400-01	.500-01	.660-00
37 86	SR 5	.186+02	.450+02	.400-02	.300-02	.400-02	.660-00
37 86	SR 6						.660+00
37 86	SR14	.186+02	.600+02	.450-00	.340-00	.450-00	.700-00
37 86	SR17	.186+02	.600+02	.300-02	.230-02	.300-02	.650-00
37 87	SR 1	.180+14	.450+02	.100+01	.750-00	.100+01	.900-01
37 87	SR 4	.180+14	.630+02	.500-01	.400-01	.500-01	.900-01
37 87	SR 5	.180+14	.450+02	.400-02	.300-02	.400-02	.900-01
37 87	SR 8						.900-01
37 87	SR14	.180+14	.600+02	.450-00	.340-00	.450-00	.900-01
37 87	SR17	.180+14	.600+02	.300-02	.230-02	.300-02	.900-01
38 85	SR 1	.650+02	.130+05	.300-00	.400-00	.100+01	.330-00
38 85	SR 6	.650+02	.160+05	.900-01	.120+00	.175-00	.910-01
38 85	SR 8						.160+00
38 85	SR 1	.490-01	.130+05	.300-00	.400-00	.100+01	.470-00
38 85	SR 6	.490-01	.180+05	.900-01	.120+00	.175-00	.130-00
38 85	SR 8						.150+00.17+00.23+00
38 89	SR 1	.505+02	.130+05	.300-00	.400-00	.100+01	.550-00
38 89	SR 6	.505+02	.180+05	.900-01	.120+00	.175-00	.280+01
38 89	SR 8						.550+00
38 89	SR25	.505+02	.300-00	.100+01	.625-00	.000	.550-00
38 89	SR21	.505+02	.300-00	.100+01	.625-00	.000	.550-00
38 89	SR22	.505+02	.300-00	.100+01	.625-00	.000	.550-00
38 89	SR23	.505+02	.300-00	.100+01	.625-00	.000	.550-00
38 90	SR 1	.150+05	.130+05	.300-00	.400-00	.100+01	.110+01
38 90	SR 6	.160+05	.180+05	.900-01	.120+00	.750-01	.550+01
38 90	SR 8						.452-00.11+01.11+01
38 91	SR 1	.400-00	.130+05	.300-00	.400-00	.100+01	.240+01
38 91	SR 6	.400-00	.160+05	.900-01	.120+00	.175-00	.740+01
38 91	SR 8						.115+01.14+01.17+01
38 92	SR 1	.110+00	.130+05	.300-00	.400-00	.100+01	.260+01
38 92	SR 6	.110+00	.180+05	.900-01	.120+00	.175-00	.800+01
38 92	SR 8						.175+01.19+01.19+01
39 90	Y 1	.268+01	.140+05	.100-03	.250-00	.100+01	.890-00
39 90	Y 6	.268+01	.180+05	.750-04	.190-00	.750-00	.440+01
39 90	Y 8						.890+00
39 91	Y 1	.580+02	.140+05	.100-03	.250-00	.100+01	.590-00
39 91	Y 6	.580+02	.180+05	.750-04	.190-00	.750-00	.290+01
39 91	Y 8						.590+00
39 911	Y 1	.350-01	.140+05	.100-03	.250-00	.100+01	.930-00
39 911	Y 6	.350-01	.180+05	.750-04	.190-00	.750-00	.300+01
39 911	Y 8						.160-00.42-00.68-00
39 92	Y 1	.150-00	.140+05	.100-03	.250-00	.100+01	.160+01
39 92	Y 6	.150-00	.180+05	.750-04	.190-00	.750-00	.590+01
39 92	Y 8						.150+01
39 93	Y 1	.420-00	.140+05	.100-03	.250-00	.100+01	.170+01
39 93	Y 6	.420-00	.180+05	.750-04	.190-00	.750-00	.650+01
39 93	Y 8						.150+01.15+01.15+01
40 93	ZR 1	.480+09	.450+03	.100-03	.250-00	.100+01	.240-01
40 93	ZR 3	.480+09	.900+03	.200-05	.500-02	.200-01	.240-01
40 93	ZR 4	.480+09	.320+03	.700-05	.200-01	.700-01	.250-01
40 93	ZR 5	.480+09	.900+03	.600-06	.150-02	.600-02	.250-01
40 93	ZR 6	.480+09	.100+04	.360-04	.900-01	.360-00	.110+00
40 93	ZR 8						.190-01.20-01.24-01
40 95	ZR 1	.633+02	.450+03	.100+03	.250-00	.100+01	.110+01
40 95	ZR 3	.633+02	.900+03	.200-05	.500-02	.200-01	.460-00
40 95	ZR 4	.633+02	.320+03	.700-05	.200-01	.700-01	.570-00
40 95	ZR 5	.633+02	.900+03	.600-06	.150-02	.600-02	.460-00
40 95	ZR 6	.633+02	.100+04	.360-04	.900-01	.360-00	.110+01

40 97 ZR 1	.710-00	.450+03	.100-03	.250-00	.100+01	.210+01	
40 97 ZR 3	.710-00	.900+03	.200-05	.500-02	.200-01	.150+01	
40 97 ZR 4	.710-00	.320+03	.700-05	.200-01	.700-01	.160+01	
40 97 ZR 5	.710-00	.900+03	.600-06	.150-02	.600-02	.150+01	
40 97 ZR 6	.710-00	.100+04	.360-04	.900-01	.360-00	.620+01	
40 97 ZR 8						.155+01	.16+01.16+01
41 931NB 1	.370+04	.760+03	.100-03	.250-00	.100+01	.380-01	
41 931NB 3	.370+04	.760+03	.200-05	.500-02	.200-01	.380-01	
41 931NB 4	.370+04	.845+03	.900-05	.200-01	.900-01	.380-01	
41 931NB 5	.370+04	.950+02	.800-06	.200-02	.800-02	.380-01	
41 931NB 6	.370+04	.100+04	.380-04	.100+00	.380-00	.120+00	
41 931NB 8						.380-01	
41 95 NB 1	.350+02	.760+03	.100-03	.250-00	.100+01	.510-00	
41 95 NB 3	.350+02	.760+03	.200-05	.500-02	.200-01	.200-00	
41 95 NB 4	.350+02	.840+03	.900-05	.200-01	.900-01	.260-00	
41 95 NB 5	.350+02	.950+02	.800-06	.200-02	.800-02	.200-00	
41 95 NB 6	.350+02	.100+04	.380-04	.100+00	.380-00	.370-00	
41 95 NB 8						.260+00	
41 97 NB 1	.510-01	.760+03	.100-03	.250-00	.100+01	.870-00	
41 97 NB 3	.510-01	.760+03	.200-05	.500-02	.200-01	.600-00	
41 97 NB 4	.510-01	.840+03	.900-05	.200-01	.900-01	.540-00	
41 97 NB 5	.510-01	.950+02	.800-06	.200-02	.800-02	.600-00	
41 97 NB 6	.510-01	.100+04	.380-04	.100+00	.380-00	.240+01	
41 97 NB 8						.640+00	
42 99 MO 1	.270+01	.500+01	.800-00	.650-00	.100+01	.510-00	
42 99 MO 3	.270+01	.300+01	.800-01	.500-01	.800-01	.480-00	
42 99 MO 4	.270+01	.450+02	.800-01	.650-01	.100+00	.480-00	
42 99 MO 8						.470-00	.47-00.47-00
43 96 TC 1	.430+01	.100+01	.500-00	.500-00	.100+01	.140+01	
43 96 TC 3	.430+01	.200+02	.500-02	.500-02	.100-01	.470-00	
43 96 TC 4	.430+01	.300+02	.150-02	.150-02	.300-02	.640-00	
43 96 TC 6	.430+01	.250+02	.100-02	.100-02	.200-02	.350-00	
43 96 TC 8	.430+01	.500+01	.450-03	.450-03	.900-03	.640-00	
43 96 TC12	.430+01	.100+02	.500-02	.500-02	.100-01	.330-02	
43 961TC 1	.360-01	.100+01	.500-00	.500-00	.100+01	.300-00	
43 961TC 3	.360-01	.200+02	.500-02	.500-02	.100-01	.420-00	
43 961TC 4	.360-01	.300+02	.150-02	.150-02	.300-02	.600-00	
43 961TC 6	.360-01	.250+02	.100-02	.100-02	.200-02	.390-00	
43 961TC 8						.155-00	.62-00.67-00
43 961TC12	.360-01	.100+02	.500-02	.500-02	.100-01	.210-01	
43 97 TC 1	.370+07	.100+01	.500-00	.500-00	.100+01	.200-01	
43 97 TC 3	.370+07	.200+02	.500-02	.500-02	.100-01	.200-01	
43 97 TC 4	.370+07	.300+02	.150-02	.150-02	.300-02	.200-01	
43 97 TC 6	.370+07	.250+02	.100-02	.100-02	.200-02	.190-01	
43 97 TC 8	.370+07	.500+01	.450-03	.450-03	.900-03	.200-01	
43 97 TC12	.370+07	.100+02	.500-02	.500-02	.100-01	.110-02	
43 971TC 1	.920+02	.100+01	.500-00	.500-00	.100+01	.900-01	
43 971TC 3	.920+02	.200+02	.500-02	.500-02	.100-01	.900-01	
43 971TC 4	.920+02	.300+02	.150-02	.150-02	.300-02	.900-01	
43 971TC 6	.920+02	.250+02	.100-02	.100-02	.200-02	.370-00	
43 971TC 8						.900-01	.90-01.90-01
43 971TC12	.920+02	.100+02	.500-02	.500-02	.100-01	.710-01	
43 99 TC 1	.730+08	.100+01	.500-00	.500-00	.100+01	.940-01	
43 99 TC 3	.730+08	.200+02	.500-02	.500-02	.100-01	.940-01	
43 99 TC 4	.730+08	.300+02	.150-02	.150-02	.300-02	.940-01	
43 99 TC 6	.730+08	.250+02	.100-02	.100-02	.200-02	.940-01	
43 99 TC 8	.730+08	.500+01	.450-03	.450-03	.900-03	.940-01	
43 99 TC12	.730+08	.100+02	.500-02	.500-02	.100-01	.470-00	
44 97 FU 1	.280+01	.730+01	.300-01	.270-00	.100+01	.150-00	
44 97 FU 3	.280+01	.250+01	.600-02	.500-01	.200-00	.770-01	
44 97 FU 6	.280+01	.160+02	.240-02	.200-01	.800-01	.130-00	
44 97 FU 8						.790-01	.11+00.15-00
44103 FU 1	.410+02	.730+01	.300-01	.270-00	.100+01	.440-00	
44103 FU 3	.410+02	.250+01	.600-02	.500-01	.200-00	.220-00	
44103 FU 6	.410+02	.160+02	.240-02	.200-01	.800-01	.620-00	

44105	FU	1	.190-00	.730+01	.300-01	.270-00	.100+01	.120+01	
44105	FU	3	.190-00	.250+01	.600-02	.500-01	.200-00	.840-00	
44105	FU	6	.190-00	.160+02	.240-02	.200-01	.800-01	.350+01	
44105	FU	8						.795-00	90-00.91-00
44106	FU	1	.365+03	.730+01	.300-01	.270-00	.100+01	.140+01	
44106	FU	3	.365+03	.250+01	.600-02	.500-01	.200-00	.130+01	
44106	FU	6	.365+03	.160+02	.240-02	.200-01	.800-01	.650+01	
44106	FU	8						.140+01	
44106	FU20		.365+03	.300-01	.100+01	.625-00	.000	.140+01	
44106	FU21		.365+03	.300-01	.100+01	.625-00	.000	.140+01	
44106	FU22		.365+03	.300-01	.100+01	.625-00	.000	.130+01	
44106	FU23		.365+03	.300-01	.100+01	.625-00	.000	.130+01	
451031FH		1	.380-01	.104+02	.200-00	.350-00	.100+01	.550-01	
451031FH		3	.380-01	.280+02	.600-02	.100-01	.300-01	.540-01	
451031FH		4	.380-01	.182+02	.600-02	.140-01	.400-01	.550-01	
451031FH		5	.380-01	.200+02	.200-02	.350-02	.100-01	.540-01	
451031FH		6	.380-01	.166+02	.100-01	.200-01	.500-01	.190-00	
451031FH		8						.550-01	
45105	FH	1	.152+01	.104+02	.200-00	.350-00	.100+01	.200-00	
45105	FH	3	.152+01	.280+02	.600-02	.100-01	.300-01	.190-00	
45105	FH	4	.152+01	.182+02	.600-02	.140-01	.400-01	.190-00	
45105	FH	5	.152+01	.200+02	.200-02	.350-02	.100-01	.190-00	
45105	FH	6	.152+01	.166+02	.100-01	.200-01	.500-01	.350-00	
45105	FH	8						.190-01	
46103	FD	1	.170+02	.500+01	.200-00	.350-00	.100+01	.640-01	
46103	FD	3	.170+02	.300+02	.200-01	.300-01	.800-01	.610-01	
46103	FD	4	.170+02	.190+02	.200-01	.300-01	.900-01	.630-01	
46103	FD	5	.170+02	.150+02	.200-02	.350-02	.100-01	.610-01	
46103	FD	8						.625-01	64-01.64-01
46109	FD	1	.570-00	.500+01	.200-00	.350-00	.100+01	.420-00	
46109	FD	3	.570-00	.300+02	.200-01	.300-01	.800-01	.420-00	
46109	FD	4	.570-00	.190+02	.200-01	.300-01	.900-01	.420-00	
46109	FD	5	.570-00	.150+02	.200-02	.350-02	.100-01	.420-00	
46109	FD	8						.420+00	
47105	AG	1	.400+02	.500+01	.100-01	.260-00	.100+01	.630-00	
47105	AG	3	.400+02	.100+02	.200-03	.500-02	.200-01	.220-00	
47105	AG	4	.400+02	.150+02	.300-03	.770-02	.300-01	.290-00	
47105	AG	6	.400+02	.300+02	.500-03	.130-01	.500-01	.160-00	
47105	AG	8						.290+00	
471101AG		1	.270+03	.500+01	.100-01	.260-00	.100+01	.170+01	
471101AG		3	.270+03	.100+02	.200-03	.500-02	.200-01	.650-00	
471101AG		4	.270+03	.150+02	.300-03	.770-02	.300-01	.840-00	
471101AG		6	.270+03	.300+02	.500-03	.130-01	.500-01	.110+01	
471101AG		8						.840+00	
47111	AG	1	.750+01	.500+01	.100-01	.260-00	.100+01	.400-00	
47111	AG	3	.750+01	.100+02	.200-03	.500-02	.200-01	.370-00	
47111	AG	4	.750+01	.150+02	.300-03	.770-02	.300-01	.380-00	
47111	AG	6	.750+01	.300+02	.500-03	.130-01	.500-01	.180+01	
47111	AG	8						.380+00	
48109	CO	1	.475+03	.200+03	.250-02	.250-00	.100+01	.110+00	
48109	CO	3	.475+03	.300+03	.250-03	.250-01	.100+00	.980-01	
48109	CO	4	.475+03	.200+03	.190-02	.190-00	.750-00	.100+00	
48109	CO	8						.100+00	
48115	CO	1	.220+01	.200+03	.250-02	.250-00	.100+01	.710-00	
48115	CO	3	.220+01	.300+03	.250-03	.250-01	.100+00	.560-00	
48115	CO	4	.220+01	.200+03	.190-02	.190-00	.750-00	.580-00	
48115	CO	8						.650-13	38-16.39-15
481151CO		1	.430+02	.200+03	.250-02	.250-00	.100+01	.610-00	
481151CO		3	.430+02	.300+03	.250-03	.250-01	.100+00	.610-00	
481151CO		4	.430+02	.200+03	.190-02	.190-00	.750-00	.610-00	
481151CO		8						.610-00	61-00.61-00
491131IN		1	.730-01	.400+02	.200-02	.250-00	.100+01	.290-00	
491131IN		3	.730-01	.600+02	.800-04	.100-01	.400-01	.190-00	
491131IN		4	.730-01	.500+02	.200-03	.400-01	.140-00	.210-00	
491131IN		5	.730-01	.400+02	.400-04	.500-02	.200-01	.190-00	

491131IN 6											.210+00
491131IN12	.730-01	.670+02	.360-03	.500-01	.180-00	.130-00					
491131IN16	.730-01	.640+01	.300-06	.100-03	.400-03	.160-00					
491141IN 1	.490+02	.480+02	.200-02	.250-00	.100+01	.970-00					
491141IN 3	.490+02	.500+02	.800-04	.100-01	.400-01	.930-00					
491141IN 4	.490+02	.580+02	.280-03	.400-01	.140-00	.940-00					
491141IN 5	.490+02	.480+02	.400-04	.500-02	.200-01	.930-00					
491141IN 6	.490+02	.570+02	.340-03	.400-01	.170-00	.450+01					
491141IN 8						.940+00					
491141IN12	.490+02	.670+02	.360-03	.500-01	.180-00	.900-00					
491141IN16	.490+02	.840+01	.800-06	.100-03	.400-03	.920-00					
49115 IN 1	.220+18	.480+02	.200-02	.250-00	.100+01	.170-00					
49115 IN 3	.220+18	.600+02	.800-04	.100-01	.400-01	.170-00					
49115 IN 4	.220+18	.580+02	.280-03	.400-01	.140-00	.170-00					
49115 IN 5	.220+18	.480+02	.400-04	.500-02	.200-01	.170-00					
49115 IN 6	.220+18	.570+02	.340-03	.400-01	.170-00	.850-00					
49115 IN 8						.170+00					
49115 IN12	.220+18	.670+02	.360-03	.500-01	.180-00	.170-00					
49115 IN16	.220+18	.840+01	.800-06	.100-03	.400-03	.170-00					
491151IN 1	.190-00	.480+02	.200-02	.250-00	.100+01	.260-00					
491151IN 3	.190-00	.600+02	.800-04	.100-01	.400-01	.190-00					
491151IN 4	.190-00	.580+02	.280-03	.400-01	.140-00	.260-00					
491151IN 5	.190-00	.480+02	.400-04	.500-02	.200-01	.190-00					
491151IN 6	.190-00	.570+02	.340-03	.400-01	.170-00	.740-00					
491151IN 8						.773-18					39-16.39-15
491151IN12	.190-00	.670+02	.360-03	.500-01	.180-00	.140-00					
491151IN16	.190-00	.840+01	.800-06	.100-03	.400-03	.160-00					
50113 SN 1	.112+03	.350+02	.500-01	.280-00	.100+01	.320-00					
50113 SN 4	.112+03	.700+02	.500-03	.280-02	.100-01	.230-00					
50113 SN 6	.112+03	.100+03	.200-01	.800-01	.300-01	.700-00					
50113 SN 8						.220-00					23-00.23-00
50113 SN15	.112+03	.350+02	.800-04	.440-03	.160-02	.160-00					
50113 SN16	.112+03	.700+02	.500-05	.280-04	.100-03	.160-00					
50125 SN 1	.950+01	.350+02	.500-01	.280-00	.100+01	.940-00					
50125 SN 4	.950+01	.700+02	.500-03	.280-02	.100-01	.940-00					
50125 SN 6	.950+01	.100+03	.200-01	.500-02	.300-00	.480+01					
50125 SN 8						.930-00					94-00.10+01
50125 SN15	.950+01	.350+02	.800-04	.440-03	.160-02	.940-00					
50125 SN16	.950+01	.700+02	.500-05	.280-04	.100-03	.930-00					
51122 SB 1	.280+01	.380+02	.300-01	.270-00	.100+01	.820-00					
51122 SB 4	.280+01	.380+02	.800-04	.500-03	.200-02	.670-00					
51122 SB 6	.280+01	.100+03	.300-02	.300-01	.100-01	.270+01					
51122 SB 8	.280+01	.100+03	.900-03	.800-02	.300-01	.670-00					
51122 SB16	.280+01	.480+01	.900-06	.800-05	.300-04	.590-00					
51124 SB 1	.600+02	.380+02	.300-01	.270-00	.100+01	.160+01					
51124 SB 4	.600+02	.380+02	.800-04	.500-03	.200-02	.920-00					
51124 SB 6	.600+02	.100+03	.300-02	.300-01	.100+00	.230+01					
51124 SB 8	.600+02	.100+03	.900-03	.800-02	.300-01	.920-00					
51124 SB16	.600+02	.480+01	.900-06	.800-05	.300-04	.570-00					
51125 SB 1	.877+03	.380+02	.300-01	.270-00	.100+01	.340-00					
51125 SB 4	.877+03	.380+02	.800+06	.500-03	.200-02	.230-00					
51125 SB 6	.877+03	.100+03	.300-02	.300-01	.100+00	.690-00					
51125 SB 8						.211-00					23-00.26-00
51125 SB16	.877+03	.480+01	.900-06	.800-05	.300-04	.150-00					
521251TE 1	.580+02	.150+02	.250-00	.360-00	.100+01	.150-00					
521251TE 3	.580+02	.300+02	.200-01	.300-01	.700-01	.140-00					
521251TE 4	.580+02	.300+02	.100-01	.200-01	.500-01	.140-00					
521251TE 5	.580+02	.300+02	.250-02	.380-02	.100-01	.140-00					
521251TE 6	.580+02	.300+02	.230-01	.340-01	.900-01	.510-00					
521251TE 8						.140+00					
521251TE10	.580+02	.300+02	.750-03	.110-02	.300-02	.110+00					
521251TE16	.580+02	.300+01	.250-03	.380-03	.360-00	.110+00					
52127 TE 1	.390-00	.150+02	.250-00	.380-00	.130+01	.240-00					
52127 TE 3	.390-00	.300+02	.200-01	.300-01	.700-01	.240-00					
52127 TE 4	.390-00	.300+02	.100-01	.200-01	.500-01	.240-00					

52127 TE 6	.390-00	.390+02	.230-01	.340-01	.900-01	.120+01	
52127 TE 8						.240+00	
52127 TE10	.390-00	.390+02	.750-03	.110-02	.300-02	.240-00	
52127 TE16	.390-00	.390+01	.250-03	.380-03	.300-00	.240-00	
521271TE 1	.105+03	.150+02	.250-00	.380-00	.100+01	.320-00	
521271TE 3	.105+03	.300+02	.200-01	.300-01	.700-01	.320-00	
521271TE 4	.105+03	.300+02	.100-01	.200-01	.500-01	.320-00	
521271TE 5	.105+03	.300+02	.250-02	.380-02	.100-01	.320-00	
521271TE 6	.105+03	.300+02	.230-01	.340-01	.900-01	.150+01	
521271TE 8						.256-00	.32-00.32-00
521271TE10	.105+03	.300+02	.750-03	.110-02	.300-02	.310-00	
521271TE16	.105+03	.900+01	.250-03	.380-03	.300-00	.300-00	
52129 TE 1	.510-01	.150+02	.250-00	.380-00	.100+01	.980-00	
52129 TE 3	.510-01	.300+02	.200-01	.300-01	.700-01	.680-00	
52129 TE 4	.510-01	.300+02	.100-01	.200-01	.500-01	.730-00	
52129 TE 5	.510-01	.300+02	.250-02	.380-02	.100-01	.680-00	
52129 TE 6	.510-01	.300+02	.230-01	.340-01	.900-01	.280+01	
52129 TE 8						.730-00	.73-00.73-00
52129 TE10	.510-01	.300+02	.750-03	.110-02	.300-02	.600-00	
52129 TE16	.510-01	.900+01	.250-03	.380-03	.300-00	.600-00	
521291TE 1	.330+02	.150+02	.250-00	.380-00	.100+01	.110+01	
521291TE 3	.330+02	.300+02	.200-01	.300-01	.700-01	.780-00	
521291TE 4	.330+02	.300+02	.100-01	.200-01	.500-01	.830-00	
521291TE 5	.330+02	.300+02	.250-02	.380-02	.100-01	.780-00	
521291TE 6	.330+02	.300+02	.230-01	.340-01	.900-01	.320+01	
521291TE 8						.790-00	.83-00.83-00
521291TE10	.330+02	.300+02	.750-03	.110-02	.300-02	.690-00	
521291TE16	.330+02	.900+01	.250-03	.380-03	.300-00	.680-00	
521311TE 1	.125+01	.150+02	.250-00	.380-00	.100+01	.160+01	
521311TE 3	.125+01	.300+02	.200-01	.300-01	.700-01	.810-00	
521311TE 4	.125+01	.300+02	.100-01	.200-01	.500-01	.970-00	
521311TE 5	.125+01	.300+02	.250-02	.380-02	.100-01	.860-00	
521311TE 6	.125+01	.300+02	.230-01	.340-01	.900-01	.260+01	
521311TE 8						.763-00	.99-00.10+01
521311TE10	.125+01	.900+01	.250-03	.380-03	.300-00	.690-00	
52132 TE 1	.320+01	.150+02	.250-00	.380-00	.100+01	.190+01	
52132 TE 3	.320+01	.300+02	.200-01	.300-01	.700-01	.960-00	
52132 TE 4	.320+01	.300+02	.100-01	.200-01	.500-01	.110+01	
52132 TE 5	.320+01	.300+02	.250-02	.380-02	.100-01	.960-00	
52132 TE 6	.320+01	.300+02	.230-01	.340-01	.900-01	.310+01	
52132 TE 8						.104+01	.11+01.11+01
52132 TE10	.320+01	.300+02	.750-03	.110-02	.300-02	.730-00	
52132 TE16	.320+01	.900+01	.250-03	.380-03	.300-00	.160-00	
53126 I 1	.133+02	.130+03	.100+01	.750-00	.100+01	.230-00	
53126 I 8						.180+01	
53126 I16	.133+02	.130+03	.300-00	.230-00	.300-00	.160-00	
53129 I 1	.630+10	.130+03	.100+01	.750-00	.100+01	.390-01	
53129 I 3	.630+10	.700+01	.400-01	.300-01	.400-01	.210-00	
53129 I 4	.630+10	.700+01	.120+00	.900-01	.120+00	.460-00	
53129 I 5	.630+10	.700+01	.500-02	.380-02	.500-02	.170-00	
53129 I 6	.630+10	.140+02	.700-01	.530-01	.700-01	.960-10	
53129 I 8						.820-01	
53129 I10	.630+10	.700+01	.500-02	.380-02	.500-02	.150-00	
53129 I16	.630+10	.130+03	.300-00	.230-00	.300-00	.600-01	
53131 I 1	.305+01	.130+03	.100+01	.750-00	.100+01	.440-00	
53131 I 8						.300+00	
53131 I16	.805+01	.130+03	.300-00	.230-00	.300-00	.230-00	
53132 I 1	.970-01	.130+03	.100+01	.750-00	.100+01	.170+01	
53132 I 3	.970-01	.700+01	.400-01	.300-01	.400-01	.850-00	
53132 I 4	.970-01	.700+01	.120+00	.900-01	.120+00	.990-00	
53132 I 5	.970-01	.700+01	.500-02	.380-02	.500-02	.950-00	
53132 I 6	.970-01	.140+02	.700-01	.530-01	.700-01	.270+01	
53132 I 8						.100+00	
53132 I10	.970-01	.700+01	.500-02	.380-02	.500-02	.640-00	
53132 I16	.970-01	.130+03	.300-00	.230-00	.300-00	.650-00	

53133	I16	.370-00	.133+03	.300-00	.230-00	.300-00	.540-00
53134	I 1	.360-01	.133+03	.100+01	.750-00	.100+01	.150+01
53134	I 8						.110+01
53134	I16	.360-01	.138+03	.300-00	.230-00	.300-00	.320-00
53135	I 1	.280-00	.133+03	.100+01	.750-00	.100+01	.130+01
53135	I 8						.990-00.11+01.11+01
53135	I16	.280-00	.138+03	.300-00	.230-00	.300-00	.520-00
55131	CS 1	.100+02	.700+02	.100+01	.750-00	.100+01	.290-01
55131	CS 3	.100+02	.420+02	.100-01	.750-02	.100-01	.210-01
55131	CS 4	.100+02	.900+02	.700-01	.500-01	.700-01	.240-01
55131	CS 5	.100+02	.980+02	.500-02	.380-02	.500-02	.210-01
55131	CS 6	.100+02	.140+03	.400-01	.300-01	.400-01	.170-01
55131	CS 8	.100+02	.140+03	.300-02	.230-02	.300-02	.240-01
55131	CS14	.100+02	.140+03	.400-00	.300-00	.400-00	.290-01
55134	CS 1	.840+03	.700+02	.100+01	.750-00	.100+01	.110+01
55134	CS 3	.840+03	.420+02	.100-01	.750-02	.100-01	.460-00
55134	CS 4	.840+03	.900+02	.700-01	.500-01	.700-01	.570-00
55134	CS 5	.840+03	.980+02	.500-02	.380-02	.500-02	.460-00
55134	CS 6	.840+03	.140+03	.400-01	.300-01	.400-01	.990-00
55134	CS 8	.840+03	.140+03	.300-02	.230-02	.300-02	.570-00
55134	CS14	.840+03	.140+03	.400-00	.300-00	.400-00	.110+01
551341CS	1	.130-00	.700+02	.100+01	.750-00	.100+01	.190-00
551341CS	3	.130-00	.420+02	.100-01	.750-02	.100-01	.110+00
551341CS	4	.130-00	.900+02	.700-01	.500-01	.700-01	.190-00
551341CS	5	.130-00	.980+02	.500-02	.380-02	.500-02	.130-00
551341CS	6	.130-00	.140+03	.400-01	.300-01	.400-01	.490-00
551341CS	8						.927-01.12+00.30-00
551341CS14		.130-00	.140+03	.400-00	.300-00	.400-00	.260-00
55135	CS 1	.110+10	.700+02	.100+01	.750-00	.100+01	.660-01
55135	CS 3	.110+10	.420+02	.100-01	.750-02	.100-01	.660-01
55135	CS 4	.110+10	.900+02	.700-01	.500-01	.700-01	.660-01
55135	CS 5	.110+10	.980+02	.500-02	.380-02	.500-02	.660-01
55135	CS 6	.110+10	.140+03	.400-01	.300-01	.400-01	.330-00
55135	CS 8	.110+10	.140+03	.300-02	.230-02	.300-02	.660-01
55135	CS14	.110+10	.140+03	.400-00	.300-00	.400-00	.660-01
55136	CS 1	.130+02	.700+02	.100+01	.750-00	.100+01	.650-00
55136	CS 3	.130+02	.420+02	.100-01	.750-02	.100-01	.290-00
55136	CS 4	.130+02	.900+02	.700-01	.500-01	.700-01	.350-00
55136	CS 5	.130+02	.980+02	.500-02	.380-02	.500-02	.290-00
55136	CS 6	.130+02	.140+03	.400-01	.300-01	.400-01	.720-00
55136	CS 8	.130+02	.140+03	.300-02	.230-02	.300-02	.350-00
55136	CS14	.130+02	.140+03	.400-00	.300-00	.400-00	.650-00
55137	CS 1	.110+05	.700+02	.100+01	.750-00	.100+01	.590-00
55137	CS 3	.110+05	.420+02	.100-01	.750-02	.100-01	.370-00
55137	CS 4	.110+05	.900+02	.700-01	.500-01	.700-01	.410-00
55137	CS 5	.110+05	.980+02	.500-02	.380-02	.500-02	.370-00
55137	CS 6	.110+05	.140+03	.400-01	.300-01	.400-01	.140+01
55137	CS 8	.110+05	.140+03	.300-02	.230-02	.300-02	.410-00
55137	CS14	.110+05	.140+03	.400-00	.300-00	.400-00	.590-00
56131	EA 1	.116+02	.650+02	.500-01	.280-00	.100+01	.380-00
56131	EA 3	.116+02	.850+01	.500-05	.280-04	.100-03	.140-00
56131	EA 4	.116+02	.975+03	.300-04	.170-03	.600-03	.190-00
56131	EA 5	.116+02	.130+02	.250-05	.140-04	.500-04	.140-00
56131	EA 6	.116+02	.650+02	.350-01	.190-00	.700-00	.110+00
56131	EA 8						.172-00.19-00.19-00
56131	EA14	.116+02	.200+04	.150-03	.830-03	.300-02	.380-00
56140	EA 1	.128+02	.650+02	.500-01	.280-00	.100+01	.230+01
56140	EA 3	.128+02	.850+01	.500-05	.280-04	.100-03	.120+01
56140	EA 4	.128+02	.975+03	.300-04	.170-03	.600-03	.140+01
56140	EA 5	.128+02	.130+02	.250-05	.140-04	.500-04	.120+01
56140	EA 6	.128+02	.650+02	.350-00	.190-00	.700-00	.420+01
56140	EA 8						.751-00.14+01.14+01
56140	EA14	.128+02	.200+04	.150-03	.830-03	.300-02	.230+01
57140	LA 1	.168+01	.500+03	.100-03	.250-00	.100+01	.190+01
57140	LA 4	.168+01	.400+03	.150-04	.400-01	.150-00	.110+01

57140	LA	8									.110+01
58141	CE	1	.320+02	.563+03	.190-03	.250-00	.100+01	.210-00			
58141	CE	3	.320+02	.563+03	.200-05	.500-02	.200-01	.160-00			
58141	CE	4	.320+02	.293+03	.250-04	.600-01	.250-00	.160-00			
58141	CE	6	.320+02	.150+04	.300-04	.750-01	.300-00	.810-00			
58141	CE	8						.160+00			
58143	CE	1	.133+01	.563+03	.100-03	.250-00	.100+01	.870-00			
58143	CE	3	.133+01	.563+03	.200-05	.500-02	.200-01	.820-00			
58143	CE	4	.133+01	.293+03	.250-04	.600-01	.250-00	.850-00			
58143	CE	6	.133+01	.150+04	.300-04	.750-01	.300-00	.380+01			
58143	CE	8						.562-00	.79-00	.85-00	
58144	CE	1	.290+03	.563+03	.100-03	.250-00	.100+01	.130+01			
58144	CE	3	.290+03	.563+03	.200-05	.500-02	.200-01	.130+01			
58144	CE	4	.290+03	.293+03	.250-04	.600-01	.250-00	.130+01			
58144	CE	6	.290+03	.150+04	.300-04	.750-01	.300-00	.630+01			
58144	CE	8						.130+01	.13+01	.13+01	
59142	FR	1	.800-00	.750+03	.100-03	.250-00	.100+01	.850-00			
59142	FR	3	.800-00	.750+03	.200-05	.500-02	.200-01	.810-00			
59142	FR	4	.800-00	.375+03	.200-04	.500-01	.200-00	.810-00			
59142	FR	6	.800-00	.150+04	.400-04	.100+00	.400-00	.390+01			
59142	FR	8						.810+00			
59143	FR	1	.137+02	.750+02	.100-03	.250-00	.100+01	.320-00			
59143	FR	3	.137+02	.750+03	.200-05	.500-02	.200-01	.320-00			
59143	FR	4	.137+02	.375+03	.200-04	.500-01	.200-00	.320-00			
59143	FR	6	.137+02	.150+04	.400-04	.100+00	.400-00	.160+01			
59143	FR	8						.320+00			
60144	MD	1	.730+08	.656+03	.100-03	.250-00	.100+01	.200+02			
60144	MD	3	.730+08	.656+03	.500-05	.100-01	.500-01	.200+02			
60144	MD	4	.730+08	.131+03	.500-04	.130-00	.500-00	.200+02			
60144	MD	6	.730+08	.150+04	.350-04	.900-01	.350-00	.100+03			
60144	MD	8						.200+02			
60147	MD	1	.113+02	.656+03	.100-03	.250-00	.100+01	.400-00			
60147	MD	3	.113+02	.656+03	.500-05	.100-01	.500-01	.310-00			
60147	MD	4	.113+02	.131+03	.500-04	.130-00	.500-00	.320-00			
60147	MD	6	.113+02	.150+04	.350-04	.900-01	.350-00	.140+01			
60147	MD	8						.290-00	.29-00	.31-00	
60149	MD	1	.830-01	.656+03	.100-03	.250-00	.100+01	.110+01			
60149	MD	3	.830-01	.656+03	.500-05	.100-01	.500-01	.970-00			
60149	MD	4	.830-01	.131+03	.500-04	.130-00	.500-00	.990-00			
60149	MD	6	.830-01	.150+04	.350-04	.900-01	.350-00	.470+01			
60149	MD	8						.688-00	.97-00	.99-00	
61147	FM	1	.920+03	.656+03	.100-03	.250-00	.100+01	.690-01			
61147	FM	3	.920+03	.656+03	.200-05	.500-02	.200-01	.690-01			
61147	FM	4	.920+03	.656+03	.600-05	.200-01	.600-01	.690-01			
61147	FM	6	.920+03	.150+04	.350-04	.900-01	.350-00	.350-00			
61147	FM	8						.690-01	.69-01	.69-01	
61149	FM	1	.220+01	.656+03	.100-03	.250-00	.100+01	.540-00			
61149	FM	3	.220+01	.656+03	.200-05	.500-02	.200-01	.420-00			
61149	FM	4	.220+01	.656+03	.600-05	.200-01	.600-01	.440-00			
61149	FM	6	.220+01	.150+04	.350-04	.900-01	.350-00	.190+01			
61149	FM	8						.440+00			
62147	SM	1	.480+14	.656+03	.100-03	.250-00	.100+01	.230+02			
62147	SM	3	.480+14	.656+03	.200-05	.500-02	.200-01	.230+02			
62147	SM	4	.480+14	.187+03	.350-04	.900-01	.350-00	.230+02			
62147	SM	6	.480+14	.150+04	.350-04	.900-01	.350-00	.115+03			
62147	SM	8						.230+02			
62151	SM	1	.370+05	.656+03	.100-03	.250-00	.100+01	.420-01			
62151	SM	3	.370+05	.656+03	.200-05	.500-02	.200-01	.420-01			
62151	SM	4	.370+05	.187+03	.350-04	.900-01	.350-00	.420-01			
62151	SM	6	.370+05	.150+04	.350-04	.900-01	.350-00	.130-00			
62151	SM	8						.420-01			
62153	SM	1	.196+01	.656+03	.100-03	.250-00	.100+01	.300-00			
62153	SM	3	.196+01	.656+03	.200-05	.500-02	.200-01	.250-00			
62153	SM	4	.196+01	.187+03	.350-04	.900-01	.350-00	.260-00			
62153	SM	6	.196+01	.150+04	.350-04	.900-01	.350-00	.110+01			

63152	EU	1	.470+04	.635+03	.100-03	.250-00	.100+01	.660-00
63152	EU	3	.470+04	.148+04	.300-05	.750-02	.300-01	.250-00
63152	EU	4	.470+04	.127+03	.250-04	.600-01	.250-00	.330-00
63152	EU	6	.470+04	.150+04	.360-04	.900-01	.360-00	.450-00
63152	EU	8						.330+00
63154	EU	1	.580+04	.635+03	.100-03	.250-00	.100+01	.130+01
63154	EU	3	.580+04	.143+04	.300-05	.750-02	.300-01	.760-00
63154	EU	4	.580+04	.127+03	.250-04	.600-01	.250-00	.860-00
63154	EU	6	.580+04	.150+04	.360-04	.900-01	.360-00	.270+01
63154	EU	8						.960+00
63155	EU	1	.621+03	.635+03	.100-03	.250-00	.100+01	.160-00
63155	EU	3	.621+03	.149+04	.300-05	.750-02	.300-01	.830-01
63155	EU	4	.621+03	.127+03	.250-04	.600-01	.250-00	.350-01
63155	EU	6	.621+03	.150+04	.360-04	.900-01	.360-00	.280-00
63155	EU	8						.950-01
64153	GO	1	.236+03	.550+03	.100-03	.250-00	.100+01	.170-00
64153	GO	4	.236+03	.460+03	.120-04	.300-01	.120+00	.990-01
64153	GO	6	.236+03	.107+04	.450-04	.110+00	.450-00	.230-00
64153	GO	8						.990-01
64159	GO	1	.750-00	.550+03	.100-03	.250-00	.100+01	.360-00
64159	GO	4	.750-00	.460+03	.120-04	.300-01	.120+00	.330-00
64159	GO	6	.750-00	.107+04	.450-04	.110+00	.450-00	.750-00
64159	GO	8						.330+00
65160	TS	1	.730+02	.670+03	.100-03	.250-00	.100+01	.350-00
65160	TS	3	.730+02	.700+03	.300-05	.750-02	.300-01	.400-00
65160	TS	6	.730+02	.100+04	.600-04	.150-00	.600-00	.110+01
65160	TS	8						.480+00
66165	EY	1	.970-01	.700+03	.100-03	.250-00	.100+01	.510-00
66165	EY	4	.970-01	.500+03	.600-05	.200-01	.600-01	.390-00
66165	EY	6	.970-01	.100+04	.600-04	.150-00	.600-00	.150+01
66165	EY	8						.390+00
66166	EY	1	.340+01	.700+03	.100-03	.250-00	.100+01	.790-00
66166	EY	4	.340+01	.500+03	.600-05	.200-01	.600-01	.780-00
66166	EY	6	.340+01	.100+03	.600-04	.150-00	.600-00	.390+01
66166	EY	8						.423-00.77-00.76-00
67166	HO	1	.110+01	.750+03	.100-03	.250-00	.100+01	.700-00
67166	HO	3	.110+01	.800+03	.200-05	.500-02	.200-01	.690-00
67166	HO	4	.110+01	.875+03	.600-05	.200-01	.600-01	.690-00
67166	HO	6	.110+01	.100+04	.640-04	.160-00	.640-00	.340+01
67166	HO	8						.690+00
68169	ER	1	.940+01	.650+03	.100-03	.250-00	.100+01	.370-00
68169	ER	3	.940+01	.650+03	.200-05	.500-02	.200-01	.190-00
68169	ER	4	.940+01	.433+03	.300-05	.750-02	.600-01	.220-00
68169	ER	6	.940+01	.100+04	.600-04	.150-00	.640-00	.580-00
68169	ER	8						.220+00
68171	ER	1	.310-00	.650+03	.100-03	.250-00	.100+01	.650-00
68171	ER	3	.310-00	.650+03	.200-05	.500-02	.200-01	.460-00
68171	ER	6	.310-00	.100+04	.600-04	.150-00	.600-00	.200+01
68171	ER	8						.490-00.49-00.50-00
69170	TM	1	.127+03	.675+03	.100-03	.250-00	.100+01	.340-00
69170	TM	3	.127+03	.335+03	.200-05	.500-02	.200-01	.340-00
69170	TM	6	.127+03	.100+04	.650-04	.160-00	.650-00	.170+01
69170	TM	8						.340+00
69171	TM	1	.694+03	.675+03	.100-03	.250-00	.100+01	.300-01
69171	TM	3	.694+03	.335+03	.200-05	.500-02	.200-01	.300-01
69171	TM	6	.694+03	.100+04	.650-04	.160-00	.650-00	.150-00
69171	TM	8						.300-01
70175	YB	1	.410+01	.665+03	.100-03	.250-00	.100+01	.160-00
70175	YB	3	.410+01	.665+03	.200-05	.130-01	.500-01	.150-00
70175	YB	6	.410+01	.100+04	.680-04	.150-00	.680-00	.710-00
70175	YB	8						.150+00
71177	LU	1	.670+01	.750+03	.100-03	.250-00	.100+01	.170-00
71177	LU	3	.670+01	.750+03	.100-05	.250-02	.100-01	.160-00
71177	LU	6	.670+01	.100+04	.680-04	.170-00	.680-00	.760-00
71177	LU	8						.160+00

72181	HF	3	.460+02	.563+03	.200-05	.500-02	.200-01	.250-00	
72181	HF	4	.460+02	.625+03	.450-04	.110+00	.450-05	.290-00	
72181	HF	5	.460+02	.350+03	.130-04	.300-01	.130-00	.250-00	
72181	HF	6	.460+02	.600+03	.150-04	.400-01	.150-00	.740-00	
72181	HF	8						.290+00	
73182	TA	1	.112+03	.240+03	.100-03	.250-00	.100+01	.110+01	
73182	TA	3	.112+03	.456+03	.390-05	.750-02	.300-01	.450-00	
73182	TA	4	.112+03	.480+03	.300-04	.800-01	.300-00	.560-00	
73182	TA	5	.112+03	.240+03	.100-05	.250-02	.100-01	.450-00	
73182	TA	6	.112+03	.300+03	.200-04	.500-01	.200-00	.100+01	
73182	TA	8						.560+00	
74181	W	1	.140+03	.100+01	.100+00	.300-00	.100+01	.200-00	
74181	W	4	.140+03	.400+01	.600-02	.200-01	.600-01	.870-01	
74181	W	6	.140+03	.900+01	.700-02	.200-01	.700-01	.470-01	
74181	W	8						.970-01	
74185	W	1	.740+02	.100+01	.100+00	.300-00	.100+01	.140-00	
74185	W	4	.740+02	.400+01	.600-02	.200-01	.600-01	.140-00	
74185	W	6	.740+02	.900+01	.700-02	.200-01	.700-01	.660-00	
74185	W	8						.140+00	
74187	W	1	.100+01	.100+01	.100+00	.300-00	.100+01	.680-00	
74187	W	4	.100+01	.400+01	.600-02	.200-01	.600-01	.440-00	
74187	W	6	.100+01	.900+01	.700-02	.200-01	.700-01	.140+01	
74187	W	8						.440-00	.44-00 .44-00
75183	RE	1	.730+02	.700+01	.500-00	.500-00	.100+01	.240-00	
75183	RE	4	.730+02	.140+02	.500-02	.500-02	.100-01	.100+00	
75183	RE	6	.730+02	.350+01	.500-02	.500-02	.100-01	.550-01	
75183	RE	8						.100+00	
75183	RE	12	.730+02	.250+02	.130-00	.130-00	.250-00	.120-02	
75183	RE	16	.730+02	.300+01	.350-02	.350-02	.700-02	.340-01	
75186	RE	1	.379+01	.700+01	.500-00	.500-00	.100+01	.380-00	
75186	RE	4	.379+01	.140+02	.500-02	.500-02	.100-01	.370-00	
75186	RE	6	.379+01	.350+01	.500-02	.500-02	.100-01	.180+01	
75186	RE	8						.370+00	
75186	RE	12	.379+01	.250+02	.130-00	.130-00	.250-00	.360-00	
75186	RE	16	.379+01	.300+01	.350-02	.350-02	.700-02	.370-00	
75187	RE	1	.180+14	.700+01	.500-00	.500-00	.100+01	.120-01	
75187	RE	4	.180+14	.140+02	.500-02	.500-02	.100-01	.120-01	
75187	RE	6	.180+14	.350+01	.500-02	.500-02	.100-01	.620-01	
75187	RE	8						.120-01	
75187	RE	12	.180+14	.250+02	.130-00	.130-00	.250-00	.120-01	
75187	RE	16	.180+14	.300+01	.350-02	.350-02	.700-02	.120-01	
75188	RE	1	.710-00	.700+01	.500-00	.500-00	.100+01	.940-00	
75188	RE	4	.710-00	.140+02	.500-02	.500-02	.100-01	.850-00	
75188	RE	6	.710-00	.350+01	.500-02	.500-02	.100-01	.390+01	
75188	RE	8						.850+00	
75188	RE	12	.710-00	.250+02	.130-00	.130-00	.250-00	.780-00	
75188	RE	16	.710-00	.300+01	.350-02	.350-02	.700-02	.800-00	
76185	CS	1	.950+02	.200+01	.100+00	.300+00	.100+01	.510-00	
76185	CS	3	.950+02	.500+01	.500-02	.200-01	.500-01	.250-00	
76185	CS	4	.950+02	.550+01	.400-02	.100-01	.400-01	.290-00	
76185	CS	6						.290+00	
76191	CS	1	.160+02	.200+01	.100+00	.300-00	.100+01	.160-00	
76191	CS	3	.160+02	.500+01	.500-02	.200-01	.500-01	.110+00	
76191	CS	4	.160+02	.950+01	.400+02	.100-01	.400-01	.120+00	
76191	CS	8						.120+00	
761911CS		1	.580-00	.200+01	.100+00	.300-00	.100+01	.600-01	
761911CS		3	.580-00	.500+01	.500-02	.200-01	.500-01	.390-01	
761911CS		4	.580-00	.550+01	.400-02	.100-01	.400-01	.490-01	
761911CS		8						.250-01	.11+00 .13-00
76193	CS	1	.131+01	.200+01	.100+00	.300-00	.100+01	.380-00	
76193	CS	3	.131+01	.500+01	.500-02	.200-01	.500-01	.380-00	
76193	CS	4	.131+01	.550+01	.400-02	.100-01	.400-01	.380-00	
76193	CS	8						.380+00	
77190	IR	1	.120+02	.200+02	.100+00	.300-00	.100+01	.370-00	

77190	IR	5	.120+02	.500+02	.200-02	.600-02	.200-01	.120+00
77190	IR	8						.160+00
77192	IR	1	.745+02	.200+02	.100+00	.300-00	.100+01	.110+01
77192	IR	3	.745+02	.500+02	.450-02	.140-01	.450-01	.500-00
77192	IR	4	.745+02	.270+02	.230-01	.700-01	.230-00	.600+00
77192	IR	5	.745+02	.500+02	.200-02	.600-02	.200-01	.500+00
77192	IR	8						.600+00
77194	IR	1	.790-00	.200+02	.100+00	.300-00	.100+01	.810-00
77194	IR	3	.790-00	.500+02	.450-02	.140-01	.450-01	.810-00
77194	IR	4	.790-00	.270+02	.230-01	.700-01	.230-00	.810-00
77194	IR	5	.790-00	.500+02	.200-02	.600-02	.200-01	.810-00
77194	IR	8						.810+00
78191	FT	1	.300+01	.240+02	.100+00	.300-00	.100+01	.700-00
78191	FT	3	.300+01	.600+02	.100-01	.300-01	.100+00	.220-00
78191	FT	4	.300+01	.200+02	.800-03	.240-02	.800-02	.310-00
78191	FT	5	.300+01	.600+02	.800-03	.240-02	.800-02	.220-00
78191	FT	8						.310+00
78193	FT	1	.180+06	.240+02	.100+00	.300-00	.100+01	.430-01
78193	FT	3	.180+06	.600+02	.100-01	.300-01	.100+00	.140-01
78193	FT	4	.180+06	.200+02	.800-03	.240-02	.800-02	.190-01
78193	FT	5	.180+06	.600+02	.800-03	.240-02	.800-02	.140-01
78193	FT	8						.190-01
781931FT	1	.350+01	.240+02	.100+00	.300-00	.100+01	.750-01	
781931FT	3	.350+01	.600+02	.100-01	.300-01	.100+00	.230-01	
781931FT	4	.350+01	.200+02	.800-03	.240-02	.800-02	.320-01	
781931FT	5	.350+01	.600+02	.800-03	.240-02	.800-02	.230-01	
781931FT	8						.320-01.32-01.32-01	
78197	FT	1	.750-00	.240+02	.100+00	.300-00	.100+01	.260-00
78197	FT	3	.750-00	.600+02	.100-01	.300-01	.100+00	.230-00
78197	FT	4	.750-00	.200+02	.800-03	.240-02	.800-02	.240-00
78197	FT	5	.750-00	.600+02	.800-03	.240-02	.800-02	.230-00
78197	FT	8						.240+00
781971FT	1	.560-01	.240+02	.100+00	.300-00	.100+01	.550-00	
781971FT	3	.560-01	.600+02	.100-01	.300-01	.100+00	.500-00	
781971FT	4	.560-01	.200+02	.800-03	.240-02	.800-02	.510-00	
781971FT	5	.560-01	.600+02	.800-03	.240-02	.800-02	.500-00	
781971FT	8						.417-00.52-00.52-00	
79196	AU	1	.560+01	.120+03	.100+00	.300-00	.100+01	.460-00
79196	AU	3	.560+01	.280+03	.300-02	.900-02	.300-01	.150-00
79196	AU	4	.560+01	.300+03	.400-02	.100-01	.400-01	.210-00
79196	AU	5	.560+01	.240+03	.500-03	.150-02	.500-02	.150-00
79196	AU	8						.210+00
79198	AU	1	.270+01	.120+03	.100+00	.300-00	.100+01	.580-00
79198	AU	3	.270+01	.280+03	.300-02	.900-02	.300-01	.410-00
79198	AU	4	.270+01	.300+03	.400-02	.100-01	.400-01	.440-00
79198	AU	5	.270+01	.240+03	.500-03	.150-02	.500-02	.410-00
79198	AU	8						.440+00
79199	AU	1	.315+01	.120+03	.100+00	.300-00	.100+01	.180-00
79199	AU	3	.315+01	.280+03	.300-02	.900-02	.300-01	.120+00
79199	AU	4	.315+01	.300+03	.400-02	.100-01	.400-01	.130-00
79199	AU	5	.315+01	.240+03	.500-03	.150-02	.500-02	.120+00
79199	AU	8						.130+00
80197	HG	1	.270+01	.100+02	.750-00	.630-00	.100+01	.970-01
80197	HG	3	.270+01	.145+02	.260-00	.220-00	.350-00	.430-01
80197	HG	4	.270+01	.135+02	.110+00	.900-01	.150-00	.520-01
80197	HG	5	.270+01	.100+02	.200-01	.100-01	.200-01	.430-01
80197	HG	8						.520-01
801971HG	1	.100+01	.100+02	.750-00	.630-00	.100+01	.300-00	
801971HG	3	.100+01	.145+02	.260-00	.220-00	.350-00	.180-00	
801971HG	4	.100+01	.135+02	.110+00	.900-01	.150-00	.190-00	
801971HG	5	.100+01	.100+02	.200-01	.100-01	.200-01	.170-00	
801971HG	8						.165-00.20-00.21-00	
80203	HG	1	.458+02	.100+02	.750-00	.630-00	.100+01	.250-00
80203	HG	3	.458+02	.145+02	.260-00	.220-00	.350-00	.150-00

80203	WG	8									.170+00
81200	TL	1	.113+01	.500+01	.450-00	.480-00	.100+01	.400-00			
81200	TL	3	.113+01	.700+01	.230-01	.240-01	.500-01	.130-00			
81201	TL	4	.113+01	.500+01	.200-01	.200-01	.400-01	.180-00			
81200	TL	6	.113+01	.700+01	.250-01	.260-01	.550-01	.350-01			
81200	TL	8	.113+01	.600+01	.270-02	.290-02	.600-02	.180-00			
81200	TL	14	.113+01	.550+01	.240-00	.260-00	.540-00	.400-00			
81201	TL	1	.300+01	.500+01	.450-00	.480-00	.100+01	.170-00			
81201	TL	3	.300+01	.700+01	.230-01	.240-01	.500-01	.110+00			
81201	TL	4	.300+01	.500+01	.200-01	.200-01	.400-01	.120+00			
81201	TL	6	.300+01	.700+01	.250-01	.260-01	.550-01	.440-00			
81201	TL	8	.300+01	.600+01	.270-02	.290-02	.600-02	.120+00			
81201	TL	14	.300+01	.550+01	.240-00	.260-00	.540-00	.170-00			
81202	TL	1	.120+02	.500+01	.450-00	.480-00	.100+01	.380-00			
81202	TL	3	.120+02	.700+01	.230-01	.240-01	.500-01	.240-00			
81202	TL	4	.120+02	.500+01	.200-01	.200-01	.400-01	.270-00			
81202	TL	6	.120+02	.700+01	.250-01	.260-01	.550-01	.940-00			
81202	TL	8	.120+02	.600+01	.270-02	.290-02	.600-02	.270-00			
81202	TL	14	.120+02	.550+01	.240-00	.260-00	.540-00	.380-00			
81204	TL	1	.110+04	.500+01	.450-00	.480-00	.100+01	.250-00			
81204	TL	3	.110+04	.700+01	.230-01	.240-01	.500-01	.250-00			
81204	TL	4	.110+04	.500+01	.200-01	.200-01	.400-01	.250-00			
81204	TL	6	.110+04	.700+01	.250-01	.260-01	.550-01	.130+01			
81204	TL	8	.110+04	.600+01	.270-02	.290-02	.600-02	.250-00			
81204	TL	14	.110+04	.550+01	.240-00	.260-00	.540-00	.250-00			
82203	FB	1	.217+01	.146+04	.800-01	.290-00	.100+01	.220-00			
82203	FB	3	.217+01	.531+03	.100-01	.400-01	.140-00	.690-01			
82203	FB	4	.217+01	.195+04	.640-02	.230-01	.800-01	.940-01			
82203	FB	6	.217+01	.365+04	.200-01	.800-01	.280-00	.510-01			
82203	FB	8						.940-01			
82210	FB	1	.710+04	.146+04	.800-01	.290-00	.100+01	.520+01			
82210	FB	3	.710+04	.531+03	.100-01	.400-01	.140-00	.100+02			
82210	FB	4	.710+04	.195+04	.640-02	.230-01	.800-01	.100+02			
82210	FB	6	.710+04	.365+04	.200-01	.800-01	.280-00	.290+02			
82210	FB	8						.157-00	.14+02	.43+02	
82212	FB	1	.440-00	.146+04	.800-01	.290-00	.100+01	.320+02			
82212	FB	3	.440-00	.531+03	.100-01	.400-01	.140-00	.810+02			
82212	FB	4	.440-00	.195+04	.640-02	.230-01	.800-01	.830+02			
82212	FB	6	.440-00	.365+04	.200-01	.800-01	.280-00	.100+02			
82212	FB	8						.790+02	.83+02	.83+02	
83206	FI	1	.640+01	.500+01	.100-01	.260-00	.100+01	.150+01			
83206	FI	3	.640+01	.600+01	.300-02	.800-01	.300-00	.590-00			
83206	FI	4	.640+01	.150+02	.150-02	.400-01	.150-00	.800-00			
83206	FI	5	.640+01	.100+02	.100-03	.260-02	.100-01	.580-00			
83206	FI	6	.640+01	.133+02	.300-03	.770-02	.300-01	.430-00			
83206	FI	8						.900+00			
83207	FI	1	.290+04	.500+01	.100-01	.260-00	.100+01	.100+01			
83207	FI	3	.290+04	.600+01	.300-02	.800-01	.300-00	.330-00			
83207	FI	4	.290+04	.150+02	.150-02	.400-01	.150-00	.450-00			
83207	FI	5	.290+04	.100+02	.100-03	.260-02	.100-01	.330-00			
83207	FI	6	.290+04	.133+02	.300-03	.770-02	.300-01	.240-00			
83207	FI	8						.450+00			
83210	FI	1	.500+01	.500+01	.100-01	.260-00	.100+01	.100+02			
83210	FI	3	.500+01	.600+01	.300-02	.800-01	.300-00	.190+02			
83210	FI	4	.500+01	.150+02	.150-02	.400-01	.150-00	.130+02			
83210	FI	5	.500+01	.100+02	.100-03	.260-02	.100-01	.170+02			
83210	FI	6	.500+01	.133+02	.300-03	.770-02	.300-01	.400+02			
83210	FI	8						.790-00	.15+02	.43+02	
83212	FI	1	.420-01	.500+01	.100-01	.260-00	.100+01	.830+02			
83212	FI	3	.420-01	.600+01	.300-02	.800-01	.300-00	.320+02			
83212	FI	4	.420-01	.150+02	.150-02	.400-01	.150-00	.930+02			
83212	FI	5	.420-01	.100+02	.100-03	.260-02	.100-01	.820+02			
83212	FI	6	.420-01	.133+02	.300-03	.770-02	.300-01	.411+03			
83212	FI	8						.930+02			

83210	BI	4	.500+01	.150+02	.150-02	.400-01	.150-00	.110+02	
83210	BI	5	.500+01	.100+02	.100-03	.260-02	.100-01	.170+02	
83210	BI	6	.500+01	.130+02	.300-03	.770-02	.300-01	.400+02	
83210	BI	8						.790-00	.15+02.43+02
83212	BI	1	.420-01	.500+01	.100-01	.260-00	.100+01	.830+02	
83212	BI	3	.420-01	.600+01	.300-02	.800-01	.300-00	.820+02	
83212	BI	4	.420-01	.150+02	.150-02	.400-01	.150-00	.830+02	
83212	BI	5	.420-01	.100+02	.100-03	.260-02	.100-01	.820+02	
83212	BI	6	.420-01	.130+02	.300-03	.770-02	.300-01	.411+03	
83212	BI	8						.830+02	
84210	PO	1	.138+03	.300+02	.600-01	.280-00	.100+01	.550+02	
84210	PO	3	.138+03	.700+02	.400-02	.200-01	.700-01	.550+02	
84210	PO	4	.138+03	.410+02	.100-01	.500-01	.170-00	.550+02	
84210	PO	5	.138+03	.600+02	.200-02	.100-01	.400-01	.550+02	
84210	PO	6	.138+03	.240+02	.600-02	.300-01	.100+00	.280+03	
84210	PO	8						.550+02	
85211	AT	1	.300-00	.270+02	.100+01	.750-00	.100+01	.610+02	
85211	AT	5	.300-00	.270+02	.300-01	.200-01	.300-01	.610+02	
85211	AT	8						.700+02	.70+02.70+02
85211	AT	11	.300-00	.135+02	.200-02	.150-02	.200-02	.610+02	
85211	AT	16	.300-00	.360+02	.300-01	.230-01	.300-01	.610+02	
88223	RA	1	.117+02	.110+04	.300-00	.400-00	.100+01	.280+03	
88223	RA	6	.117+02	.164+05	.150-00	.200-00	.500-00	.280+03	
88223	RA	8						.280+03	
88224	RA	1	.364+01	.110+04	.300-00	.400-00	.100+01	.280+03	
88224	RA	6	.364+01	.164+05	.150-00	.200-00	.500-00	.280+03	
88224	RA	8						.245+03	.28+03.28+03
88226	RA	1	.590+06	.810+04	.300-00	.400-00	.100+01	.110+03	
88226	RA	6	.590+06	.164+05	.400-01	.300-01	.100+00	.110+03	
88226	RA	8						.110+03	
88228	RA	1	.240+04	.810+04	.300-00	.400-00	.100+01	.230+03	
88228	RA	6	.240+04	.164+05	.400-01	.300-01	.100+00	.190+03	
88228	RA	8						.710-00	.22+02.14+03
89227	AC	1	.800+04	.240+05	.100-03	.250-00	.100+01	.200+03	
89227	AC	3	.800+04	.240+05	.100-05	.300-02	.100-01	.620+02	
89227	AC	4	.800+04	.240+04	.500-04	.130-00	.500-00	.620+02	
89227	AC	6	.800+04	.365+05	.300-04	.800-01	.300-00	.100+04	
89227	AC	8						.480+01	.21+03.32+03
89228	AC	1	.260-00	.240+05	.100-03	.250-00	.100+01	.230+03	
89228	AC	3	.260-00	.240+05	.100-05	.300-02	.100-01	.550+02	
89228	AC	4	.260-00	.240+04	.500-04	.250-00	.500-00	.560+02	
89228	AC	6	.260-00	.365+05	.300-04	.800-01	.300-00	.970+03	
89228	AC	8						.897-00	.22+02.14+03
90227	TH	1	.184+02	.570+05	.100-03	.250-00	.100+01	.200+03	
90227	TH	3	.184+02	.220+05	.500-05	.100-01	.500-01	.610+02	
90227	TH	4	.184+02	.570+05	.500-05	.100-01	.500-01	.610+02	
90227	TH	6	.184+02	.365+05	.700-04	.180-00	.700-00	.990+03	
90227	TH	8						.827+02	.28+03.33+03
90228	TH	1	.700+03	.570+05	.100-03	.250-00	.100+01	.230+03	
90228	TH	3	.700+03	.220+05	.500-05	.100-01	.500-01	.560+02	
90228	TH	4	.700+03	.570+05	.500-05	.100-01	.500-01	.560+02	
90228	TH	6	.700+03	.365+05	.700-04	.180-00	.700-00	.970+03	
90228	TH	8						.110+03	.31+03.33+03
90230	TH	1	.290+08	.570+05	.100-03	.250-00	.100+01	.480+02	
90230	TH	3	.290+08	.220+05	.500-05	.100-01	.500-01	.480+02	
90230	TH	4	.290+08	.570+05	.500-05	.100-01	.500-01	.480+02	
90230	TH	6	.290+08	.365+05	.700-04	.180-00	.700-00	.240+03	
90230	TH	8						.480+02	.40+02.48+02
90231	TH	1	.107+01	.570+05	.100-03	.250-00	.100+01	.180-00	
90231	TH	3	.107+01	.220+05	.500-05	.100-01	.500-01	.140-00	
90231	TH	4	.107+01	.570+05	.500-05	.100-01	.500-01	.160-00	
90231	TH	6	.107+01	.365+05	.700-04	.180-00	.700-00	.560+00	
90231	TH	8						.110+00	.11+00.11+00
90232	TH	1	.510+13	.570+05	.100-03	.250-00	.100+01	.620+02	
90232	TH	3	.510+13	.220+05	.500-05	.100-01	.500-01	.410+02	
90232	TH	4	.510+13	.570+05	.500-05	.100-01	.500-01	.410+02	
90232	TH	6	.510+13	.365+05	.700-04	.130-00	.700-00	.270+03	
90232	TH	8						.410+02	.41+02.65+02
90234	TH	1	.241+02	.570+05	.100-03	.250-00	.100+01	.910-00	
90234	TH	3	.241+02	.570+05	.500-05	.100-01	.500-01	.900-00	
90234	TH	4	.241+02	.570+05	.500-05	.100-01	.500-01	.900-00	
90234	TH	6	.241+02	.365+05	.500-05	.100-01	.700-00	.450+01	
90234	TH	8						.900+00	
91230	PA	1	.177+02	.410+05	.100-03	.250-00	.100+01	.670-01	
91230	PA	3	.177+02	.510+05	.400-05	.100-01	.400-01	.310-01	
91230	PA	6	.177+02	.365+05	.450-04	.110+00	.450-00	.110+00	
91230	PA	8						.320+01	.50+02.68+02
91231	PA	1	.130+08	.410+05	.100-03	.250-00	.100+01	.140+03	

91231	PA	3	.130+08	.510+05	.400-05	.100-01	.400-01	.790+02
91231	PA	4	.130+08	.520+05	.500-05	.130-01	.500-01	.630+02
91231	PA	6	.130+08	.365+05	.450-04	.110+00	.450-00	.750+03
91231	PA	8						.510+02.52+02.70+02
91233	PA	1	.274+02	.410+05	.100-03	.250-00	.100+01	.320-00
91233	PA	3	.274+02	.510+05	.400-05	.100-01	.400-01	.150-00
91233	PA	4	.274+02	.580+05	.500-05	.130-01	.500-01	.180-00
91233	PA	6	.274+02	.365+05	.450-04	.110+00	.450-00	.410-00
91233	PA	8						.180+00
92230	U	1	.208+02	.100+03	.100-03	.250-00	.100+01	.350+03
92230	U	3	.208+02	.150+02	.110-04	.280-01	.110+00	.350+03
92230	U	6	.208+02	.380+03	.110-04	.280-01	.110+00	.180+04
92230	U	8						.348+03.35+03.35+03
92232	U	1	.270+05	.100+03	.100-03	.250-00	.100+01	.280+03
92232	U	3	.270+05	.150+02	.110-04	.280-01	.110+00	.110+03
92232	U	6	.270+05	.300+03	.110-04	.280-01	.110+00	.120+04
92232	U	8						.552+02.76+02.19+03
92233	U	1	.590+08	.100+03	.100-03	.250-00	.100+01	.500+02
92233	U	3	.590+08	.150+02	.110-04	.280-01	.110+00	.500+02
92233	U	6	.590+08	.300+03	.110-04	.280-01	.110+00	.250+03
92233	U	8						.500+02
92234	U	1	.910+08	.100+03	.100-03	.250-00	.100+01	.490+02
92234	U	3	.910+08	.150+02	.110-04	.280-01	.110+00	.490+02
92234	U	6	.910+08	.300+03	.110-04	.280-01	.110+00	.240+03
92234	U	8						.490+02
92235	U	1	.260+12	.100+03	.100-03	.250-00	.100+01	.460+02
92235	U	3	.260+12	.150+02	.110-04	.280-01	.110+00	.460+02
92235	U	6	.260+12	.300+03	.110-04	.280-01	.110+00	.230+03
92235	U	8						.461+02.46+02.46+02
92236	U	1	.870+10	.100+03	.100-03	.250-00	.100+01	.470+02
92236	U	3	.870+10	.150+02	.110-04	.280-01	.110+00	.477+02
92236	U	6	.870+10	.300+03	.110-04	.280-01	.110+00	.230+03
92236	U	8						.470+02
92238	U	1	.160+13	.100+03	.100-03	.250-00	.100+01	.430+02
92238	U	3	.160+13	.150+02	.110-04	.280-01	.110+00	.430+02
92238	U	6	.160+13	.300+03	.110-04	.280-01	.110+00	.220+03
92238	U	8						.430+02
93237	NP	1	.800+09	.390+05	.100-03	.250-00	.100+01	.490+02
93237	NP	3	.800+09	.640+05	.300-05	.750-02	.300-01	.490+02
93237	NP	4	.800+09	.140+05	.500-05	.130-01	.500-01	.490+02
93237	NP	6	.800+09	.365+05	.540-04	.110+00	.450-00	.250+03
93237	NP	8						.490+02.49+02.49+02
93239	NP	1	.233+01	.150+05	.100-03	.250-00	.100+01	.290-00
93239	NP	3	.233+01	.640+05	.300-05	.750-02	.300-01	.210-00
93239	NP	4	.233+01	.540+05	.500-05	.130-01	.500-01	.220-00
93239	NP	6	.233+01	.365+05	.540-04	.110+00	.450-00	.980-00
93239	NP	8						.160-00.16-00.16-00
94238	PU	1	.330+05	.350+05	.300-04	.250-00	.100+01	.570+02
94238	PU	3	.330+05	.320+05	.600-06	.500-02	.200-01	.570+02
94238	PU	4	.330+05	.300+05	.450-05	.380-01	.450-00	.570+02
94238	PU	6	.330+05	.365+05	.240-04	.200-00	.450-00	.280+03
94238	PU	8	.330+05	.120+03	.300-04	.125-00	.000	.570+02
94238	PU20		.330+05	.300-04	.100+01	.625-00	.000	.550-00
94238	PU21		.330+05	.300-04	.100+01	.625-00	.000	.550-00
94238	PU22		.330+05	.300-04	.100+01	.625-00	.000	.550-00
94238	PU23		.330+05	.300-04	.100+01	.625-00	.000	.550-00
94239	PU	1	.890+07	.650+05	.300-04	.250-00	.100+01	.530+02
94239	PU	3	.890+07	.320+05	.600-06	.500-02	.200-01	.530+02
94239	PU	4	.890+07	.300+05	.450-05	.380-01	.450-00	.530+02
94239	PU	6	.890+07	.365+05	.240-04	.200-00	.450-00	.270+03
94239	PU	8	.890+07	.120+03	.300-04	.125-00	.000	.530+02
94239	PU20		.890+07	.300-04	.100+01	.625-00	.000	.520-00
94239	PU21		.890+07	.300-04	.100+01	.625-00	.000	.520-00
94239	PU22		.890+07	.300-04	.100+01	.625-00	.000	.520-00
94239	PU23		.890+07	.300-04	.100+01	.625-00	.000	.520-00
94240	PU	1	.240+07	.650+05	.300-04	.250-00	.100+01	.530+02
94240	PU	3	.240+07	.320+05	.600-06	.500-02	.200-01	.530+02
94240	PU	4	.240+07	.300+05	.450-05	.380-01	.450-00	.530+02
94240	PU	6	.240+07	.365+05	.240-04	.200-00	.450-00	.270+03
94240	PU	8	.240+07	.120+03	.300-04	.125-00	.000	.530+02
94240	PU20		.240+07	.300-04	.100+01	.625-00	.000	.520-00
94240	PU21		.240+07	.300-04	.100+01	.625-00	.000	.520-00
94240	PU22		.240+07	.300-04	.100+01	.625-00	.000	.520-00
94240	PU23		.240+07	.300-04	.100+01	.625-00	.000	.520-00
94241	PU	1	.480+04	.650+05	.300-04	.250-00	.100+01	.230+01

94240	PU 3	.240+07	.320+05	.600-06	.500-02	.200-01	.530+02	
94240	PU 4	.240+07	.300+05	.450-05	.380-01	.450-00	.530+02	
94240	PU 6	.240+07	.365+05	.240-04	.200-00	.450-00	.270+03	
94240	PU 8	.240+07	.120+03	.300-04	.125-00	.000	.530+02	
94240	PU20	.240+07	.300-04	.100+01	.625-00	.000	.520-00	
94240	PU21	.240+07	.300-04	.100+01	.625-00	.000	.520-00	
94240	PU22	.240+07	.300-04	.100+01	.625-00	.000	.520-00	
94240	PU23	.240+07	.300-04	.100+01	.625-00	.000	.520-00	
94241	PU 1	.480+04	.650+05	.300-04	.250-00	.100+01	.230+01	
94241	PU 3	.480+04	.320+05	.600-06	.500-02	.200-01	.250+01	
94241	PU 4	.480+04	.300+05	.450-05	.380-01	.450-00	.100+01	
94241	PU 6	.480+04	.365+05	.240-04	.200-00	.450-00	.140+02	
94241	PU 8						.133-01	.30-01.18-00
94241	PU20	.480+04	.300-04	.100+01	.625-00	.000	.570-00	
94241	PU21	.480+04	.300-04	.100+01	.625-00	.000	.600-00	
94241	PU22	.480+04	.300-04	.100+01	.625-00	.000	.570-00	
94241	PU23	.480+04	.300-04	.100+01	.625-00	.000	.570-00	
94242	PU 1	.140+09	.650+05	.300-04	.250-00	.100+01	.510+02	
94242	PU 3	.140+09	.320+05	.600-06	.500-02	.200-01	.510+02	
94242	PU 4	.140+09	.300+05	.450-05	.380-01	.450-00	.510+02	
94242	PU 6	.140+09	.365+05	.240-04	.200-00	.450-00	.250+03	
94242	PU 8	.140+09	.120+03	.300-04	.125-00	.000	.510+02	
94242	PU20	.140+09	.300-04	.100+01	.625-00	.000	.490-00	
94242	PU21	.140+09	.300-04	.100+01	.625-00	.000	.490-00	
94242	PU22	.140+09	.300-04	.100+01	.625-00	.000	.490-00	
94242	PU23	.140+09	.300-04	.100+01	.625-00	.000	.490-00	
95241	AM 1	.170+06	.200+05	.100-03	.250-00	.100+01	.570+02	
95241	AM 3	.170+06	.270+05	.300-05	.750-02	.300-01	.570+02	
95241	AM 4	.170+06	.348+04	.350-04	.880-01	.350-00	.570+02	
95241	AM 6	.170+06	.365+05	.250-04	.630-01	.250-00	.280+03	
95241	AM 8	.167+06	.120+03	.300-04	.125-00	.000	.570+02	
95242	AM 1	.667+00	.200+05	.100-03	.250+00	.100+01	.670+02	
95242	AM 3	.667+00	.270+05	.300-05	.750-02	.030+00	.660+02	
95242	AM 4	.667+00	.348+04	.350-04	.088+00	.350+00	.630+02	
95242	AM 6	.667+00	.365+05	.250-04	.063+00	.250+00	.340+03	
95242	AM 8						.540+00	.13+02.40+02
952421	AM 1	.560+05	.200+05	.100-03	.250+00	.100+01	.610+02	
952421	AM 3	.560+05	.270+05	.300-05	.750-02	.030+00	.610+02	
952421	AM 4	.560+05	.348+04	.350-04	.088+00	.350+00	.570+02	
952421	AM 6	.560+05	.365+05	.250-04	.063+00	.250+00	.300+03	
952421	AM 8						.340+00	.13+02.40+02
95243	AM 1	.290+07	.200+05	.100-03	.250-00	.100+01	.540+02	
95243	AM 3	.290+07	.270+05	.300-05	.750-02	.300-01	.540+02	
95243	AM 4	.290+07	.348+04	.350-04	.880-01	.350-00	.540+02	
95243	AM 6	.290+07	.365+05	.250-04	.630-01	.250-00	.270+03	
95243	AM 8						.540+02	.54+02.54+02
96242	CM 1	.163+03	.240+05	.100-03	.250-01	.100+01	.800+02	
96242	CM 3	.163+03	.240+05	.200-05	.500-02	.200-01	.780+02	
96242	CM 4	.163+03	.300+04	.400-04	.100+00	.400-00	.790+02	
96242	CM 6	.163+03	.365+05	.300-04	.750-01	.300-00	.400+03	
96242	CM 8						.630+02	.63+02.64+02
96243	CM 1	.130+05	.240+05	.100-03	.250-00	.100+01	.600+02	
96243	CM 3	.130+05	.240+05	.200-05	.500-02	.200-01	.600+02	
96243	CM 4	.130+05	.300+04	.400-04	.100+00	.400-00	.600+02	
96243	CM 6	.130+05	.365+05	.300-04	.750-01	.300-00	.300+03	
96243	CM 8						.600+02	.60+02.60+02
96244	CM 1	.670+04	.240+05	.100-03	.250-00	.100+01	.600+02	
96244	CM 3	.670+04	.240+05	.200-05	.500-02	.200-01	.600+02	
96244	CM 4	.670+04	.300+04	.400-04	.100+00	.400-00	.600+02	
96244	CM 6	.670+04	.365+05	.300-04	.750-01	.300-00	.300+03	
96244	CM 8						.600+02	.60+02.60+02
96245	CM 1	.730+07	.240+05	.100-03	.250-00	.100+01	.560+02	
96245	CM 3	.730+07	.240+05	.200-05	.500-02	.200-01	.560+02	
96245	CM 4	.730+07	.300+04	.400-04	.100+00	.400-00	.560+02	
96245	CM 6	.730+07	.365+05	.300-04	.750-01	.300-00	.280+03	

96245	CM	8									.550+02.55+02.55+02
96246	CM	1	.240+07	.240+05	.100-03	.250-00	.100+01	.560+02			
96246	CM	3	.240+07	.240+05	.200-05	.500-02	.200-01	.560+02			
96246	CM	6	.240+07	.365+05	.300-04	.750-01	.300-00	.280+03			
96246	CM	8						.560+02			
97249	BK	1	.290+03	.650+05	.300-04	.250-00	.100+01	.380+01			
97249	BK	6	.290+03	.365+05	.240-04	.200-00	.800-00	.200+02			
97249	BK	8						.264-01.44-01.20-00			
98249	CF	1	.170+06	.650+05	.300-04	.250-00	.100+01	.600+02			
98249	CF	6	.170+06	.365+05	.240-04	.200-00	.800-00	.300+03			
98249	CF	8						.600+02			
98250	CF	1	.370+04	.650+05	.300-04	.250-00	.100+01	.620+02			
98250	CF	6	.370+04	.365+05	.240-04	.200-00	.800-00	.310+03			
98250	CF	8						.620+02.62+02.62+02			
98252	CF	1	.804+03	.650+05	.300-04	.250-00	.100+01	.210+03			
98252	CF	6	.804+03	.365+05	.240-04	.200-00	.800-00	.110+04			
98252	CF	8						.210+03			

APPENDIX E

PROGRAM LISTING

APPENDIX E

PROGRAM LISTING

The order of the program listing shown here is:

MAIN

LIBB

LIBK

DEP

ERFXP

SIMPS

PASSIG

CHAI NT

ORDOSE

ORDOT1

ORDOT2

QNT1

QLUNT1

QLMIT1

UNIT

PROGRAM LISTING

PROGRAM MAIN 74/74 OPT=C TRACE FTN 4.2+P380 12/16/74 14.40.20.

	PROGRAM MAIN(INPUT,OUTPUT,TAPE1=INPUT,TAPE2=OUTPUT,FUNCH,	MAIN	2
	ORGLIB,RIBLIB,TAPE26=ORGLIB,TAPE29=RIBLIB,TAPE1)	9/9/74	9
	COMMON HFP,NAC,NTOT,NOFNUC(96),SKIP(96),MASS(600,2),LAMBDA(600),	COMN	2
	1 CHAINS,DKFECT(450,3),REC(600),ML(600),ICL(600)	COMN	3
5	COMMON MA(600),LAMBIO(1200),FWAT(1200),FAIR(1200),F2P(1200)	COMN	4
	COMMON IORG(1200),ENG(1200,3),F1(600),I8-G(600),IEND(600)	COMN	5
	COMMON COSTIM(10),D3,04,05,UPTIME(5),ISM(600)	COMN	6
	COMMON BRATE(5),R(10),X(10),SY(10),SZ(10),DIAM	COMN	7
10	COMMON SUMDOS(50,10,10),TITLE2(8),NDT,KORG(20),TLH,TLN	COMN	8
	COMMON IPAGE,ORGAN(2,10)	COMN	9
	COMMON/NAM/ORGNAH(2,25)	COMN	10
	REAL LAMBIO,LAMBDA,MASS	COMN	11
	INTEGER CHAINS,SKIP	COMN	12
15	LOGICAL FISPRD,ACTPRD	COMN	13
	COMMON ACTPRD,FISPRD,KR,KUPT,DOY	COMN	14
	COMMON LIBT(7)	9/9/74	3
	DIMENSION SHASS(600),OTLH(5),OTLN(5),ORGANS(10)	MAIN	5
	DIMENSION QUANTY(600)	MAIN	6
20	DIMENSION STUB(5),SIGZ(10,5),SIGY(10,5),EGQ(10,5)	MAIN	7
	INTEGER REC,ORGANS	MAIN	8
	REAL KSQD	MAIN	9
	DIMENSION NAME(600),M(600)	MAIN	10
	DIMENSION CTLH(10),CTLN(10),ISL(600)	MAIN	11
	DIMENSION ITA8(21)	MAIN	12
25	DIMENSION ISPACE(10),LFMT(8),MFMT(5),NFMT(6),JFMT(9),IFIELD(10)	MAIN	13
	DATA (LFMT(I),I=1,5) /"(1H," ", " ", "X,1GH", "ISTANC", "E /,7X,"/	MAIN	14
	DATA LFMT(7),LFMT(8) /"(A6,A5", ")/	MAIN	15
	DATA (MFMT(I),I=1,5) /"(1H " " " "X,5GH", "OSE TI", "NE)/	MAIN	16
30	DATA (NFMT(I),I=1,6) /"(1H," " " "X,1H", "G8.0", "7H M8T", "ERS//)	MAIN	17
	1"/	MAIN	18
	DATA (JFMT(I),I=1,6) /"(1H " " " "X,F7.1", "1X,A5", "2X,1P", "E11.2	MAIN	19
	1,"/	MAIN	20
	DATA JFMT(8),JFMT(9) /"(E11.2", ")/	MAIN	21
	DATA IFIELD/"8", "1", "2", "3", "4", "5", "6", "7", "8", "9"/	MAIN	22
35	DATA CH,OH/13.0,232./	MAIN	23
	DATA ((ORGNAM(I,J),I=1,2),J=1,23) /"TOTAL ", "BODY ", "BODY W",	MAIN	24
	1"ATER ", " KIDN", "EYS ", " LIV", "ER ", " SPLE", "EN ",	MAIN	25
	2" BON", "E ", " FA", "T ", " LUN", "GS ", " ADRLN",	MAIN	26
	3"ALS ", " TES", "TES ", " OVAR", "IES ", " SKI", "N ",	MAIN	27
40	4" BRA", "IN ", " MUSC", "LE ", " PRST", "ATE ", " THYR",	MAIN	28
	5"OID ", " PANCR", "EAS ", " HEA", "RT ", " GI TI", "SSUE ",	MAIN	29
	6" STOM", "ACH ", "SMALL ", "INTEST", "UP-LG-", "INTEST", "LO-LG-",	MAIN	30
	7"INTEST"/	MAIN	31
	NAMELIST/INPUT/NEXT,MET,AH,BH,CH,DH,KSQD,CZ,CY,EN,UBAR,D3,D4,D5,	MAIN	32
45	1H,DIAM,UPTIME,BRATE,STUB,IPR,IFU,IOP,SIGZ,SIGY,R,COSTIM,ORGANS	MAIN	33
	UNIT 29 IS FOR RIBU LIBRARY, USE LOGICAL FILE NAME AND PERMANENT	MAIN	34
	FILE NAME OF -RIBLIB- ON ATTACH CARD.	MAIN	35
	C	MAIN	36
	C	MAIN	37
50	UNIT 28 IS FOR THE ORGAN DATA LIBRARY, USE LOGICAL AND PERMANENT	MAIN	38
	FILE NAME OF -ORGLIB- ON ATTACH CARD.	MAIN	39
	C	MAIN	40
	CALL DATE(DOY)	MAIN	41
	IER=0	MAIN	42
	MET=-1	MAIN	43
55	IPAGE=1	MAIN	44
	100 READ 1050, TITLE2	MAIN	45
	READ (5,INPUT)	MAIN	45

		KASE=1	MAIN	46
		IF (NEXT.GT.3.AND.NEXT.LE.4) GO TO 110	MAIN	47
60		PRINT 900,NEXT	MAIN	48
		STOP	MAIN	49
	110	GO TO (120,130,270,810),NEXT	MAIN	50
	C	READ NUCLIDE DATA LIBRARY	MAIN	51
	120	CONTINUE	MAIN	52
65		CALL LIBK	MAIN	53
		CALL LIB8	MAIN	54
	130	DO 140 I=1,CHAINS	MAIN	55
		SKIP(I)=0	MAIN	56
	140	CONTINUE	MAIN	57
70		IEP=0	MAIN	58
	C	INITIALIZE NUCLIDE MASS ARRAY	MAIN	59
		DO 150 I=1,NTOT	MAIN	60
		MASS(I,2)=0.	MAIN	61
	150	MASS(I,1)=J.0	MAIN	62
75		N1=NFP	MAIN	63
		N2=NTOT	MAIN	64
	C	READ NUCLIDE INVENTORY FROM CARDS	MAIN	65
		READ 850, NUMBER	MAIN	66
		READ 880, (NAME(I),M(I),ISL(I),QUANTY(I),I=1,NUMBER)	MAIN	67
80		LIM=282	MAIN	68
		PRINT 855,IPAGE,DOY	MAIN	69
		PRINT 856,(LIBT(I),I=1,7)	9/9/74	4
		PRINT 860	MAIN	70
		ISTART=1	MAIN	71
85	160	IPAGE=IPAGE+1	MAIN	72
		ISTOP=MIND(NUMBER,LIM)	MAIN	73
		DO 170 I=ISTART,ISTOP,6	MAIN	74
		JSTOP=MIND(I+5,ISTOP)	MAIN	75
	170	PRINT 870, (NAME(J),M(J),ISL(J),QUANTY(J),J=I,JSTOP)	MAIN	76
90		IF (ISTOP.EQ.NUMBER) GO TO 180	MAIN	77
		ISTART=ISTOP+1	MAIN	78
		PRINT 820,IPAGE,DOY	MAIN	79
		LIM=LIM+330	MAIN	80
		GO TO 160	MAIN	81
95	C		MAIN	82
	C	TRY TO IDENTIFY NUCLIDES READ FROM CARDS	MAIN	83
	180	DO 210 I=1,NUMBER	MAIN	84
		DO 190 J=1,NTOT	MAIN	85
		IF (NAME(I).EQ.REC(J).AND.M(I).EQ.ML(J)) GO TO 200	MAIN	86
100	190	CONTINUE	MAIN	87
	C		MAIN	88
	C	IF CAN NOT IDENTIFY NUCLIDE -STOP	MAIN	89
		PRINT 890, NAME(I),M(I)	MAIN	90
		IER=IER+1	MAIN	91
105	C	CONVERT CURIES TO CURIE-SEC	MAIN	92
	200	MASS(J,1)=QUANTY(I)/LAMBDA(J)	MAIN	93
		ICL(J)=ISL(I)	MAIN	94
		IF (ISL(I).GE.1.AND.ISL(I).LE.3) GO TO 210	MAIN	95
	C	IF NO SOLVEILITY INDEX ON NUCLIDE PRINT OFFENDER	MAIN	96
110		IF (ICL(I).GE.1.AND.ICL(I).LE.3) GO TO 210	MAIN	97
		PRINT 1060, NAME(I),M(I),ICL(I)	MAIN	98
		IER=IER+1	MAIN	99
	210	CONTINUE	MAIN	100
	C	ELIMINATE UNUSED CHAINS	MAIN	101

115	C	FISPRD = FALSE IF NO FISSION PRODUCTS PRESENT	MAIN	112
	C	ACTPRD = FALSE IF NO ACTIVATION PRODUCTS PRESENT	MAIN	113
		ACTPRD=.FALSE.	MAIN	114
		FISPRD=.FALSE.	MAIN	115
		ISOTOP=0	MAIN	116
120		DO 250 K=1,CHAINS	MAIN	117
		LIM=NOFNUC(K)	MAIN	118
		IF (SKIP(K).NE.0) GO TO 240	MAIN	119
		DO 220 I=1,LIM	MAIN	110
		L=ISOTOP+I	MAIN	111
125		IF (MASS(L,1).GT.1.E-30.AND.LAMBDA(L).GT.1.E-30) GO TO 230	MAIN	112
	220	CONTINUE	MAIN	113
		SKIP(K)=LIM	MAIN	114
		GO TO 240	MAIN	115
	230	CONTINUE	MAIN	116
130		FISPRD=.TRUE.	MAIN	117
	240	ISOTOP=ISOTOP+LIM	MAIN	118
	250	CONTINUE	MAIN	119
		DO 260 L=N1,N2	MAIN	120
		IF (MASS(L,1).LT.1.E-30.OR.LAMBDA(L).LT.1.E-30) GO TO 260	MAIN	121
135		ACTPRD=.TRUE.	MAIN	122
	260	CONTINUE	MAIN	123
	C	COUNT DOSE TIMES, DISTANCES, AND UPTAKE TIMES	MAIN	124
	270	NUPT=0	MAIN	125
		NR=0	MAIN	126
140		NDT=0	MAIN	127
		IF (NEXT.LT.3) GO TO 275	MAIN	128
		PRINT 820,IPAGE,DOY	MAIN	129
		IPAGE=IPAGE + 1	MAIN	130
	275	DO 320 I=1,10	MAIN	131
145		IF (R(I)) 290,290,280	MAIN	132
	280	NR=I	MAIN	133
	290	IF (DOSTIM(I)) 310,310,300	MAIN	134
	300	NDT=I	MAIN	135
	310	IF (NUPT.GE.5.OR.I.GT.5.OR.UPTIME(I).LE.0.) GO TO 320	MAIN	136
150		NUPT=NUPT+1	MAIN	137
		UPTIME (NUPT)=UPTIME (I)	MAIN	138
	320	CONTINUE	MAIN	139
		CALL UNIT (DTLH,DTLN,NUPT,UPTIME)	MAIN	140
		IO=0	MAIN	141
155		DO 330 I=1,20	MAIN	142
	330	KORG(I)=0	MAIN	143
		DO 340 I=1,10	MAIN	144
		IND=ORGANS (I)	MAIN	145
		IF (IND.LE.0) GO TO 340	MAIN	146
160		IO=IO+1	MAIN	147
		KORG(IND)=1	MAIN	148
	340	CONTINUE	MAIN	149
		IF (IO.GT.0) GO TO 350	MAIN	150
		PRINT 910, ORGANS	MAIN	151
165		IER=IER+1	MAIN	152
	350	IF (UBAR.LT.1.E-30) IER=IER+1	MAIN	153
		IF (SRATE(1).LE.0.) IER=IER+1	MAIN	154
		IF (MET.LT.0.OR.MST.GT.10) IER=IER+1	MAIN	155
		IF (LIM.LE.220) GO TO 355	MAIN	156
170		IPAGE=IPAGE + 1	MAIN	157
		PRINT 820,IPAGE,DOY	MAIN	158

	355	PRINT 920,UBAR,BRATE,MET	MAIN	159
		IM=M	MAIN	160
		PRINT 356,IM	MAIN	161
175	356	FORMAT(1H,"M=",15," METERS")	MAIN	162
		IGC=MET	MAIN	163
		IF (MET.EQ.0) GO TO 815	MAIN	164
		IF (MET.GT.3) IGO=3	MAIN	165
		GO TO (350,330,390), IGO	MAIN	166
180	360	IF (AH.LE.0.) IER=IER+1	MAIN	167
		IF (BH.LE.0.) IER=IER+1	MAIN	168
		IF (CH.LE.0.) IER=IER+1	MAIN	169
		IF (DH.LE.0.) IER=IER+1	MAIN	170
		IF (KSQD.LE.0.) IER=IER+1	MAIN	171
185		IF (STUB(1).LE.0.) IER=IER+1	MAIN	172
		PRINT 930, AH,BH,CH,DH,KSQD,STUB(1)	MAIN	173
		IF (NUPT.EQ.1) GO TO 390	MAIN	174
		DO 370 I=2,NUPT	MAIN	175
		IF (STUB(I).LE.1.E-30) STUB(I)=STUB(I-1)	MAIN	176
190		IF (BRATE(I).LE.0.) BRATE(I)=BRATE(I-1)	MAIN	177
	370	CONTINUE	MAIN	178
		GO TO 390	MAIN	179
	380	IF (CZ.LE.0.) IER=IER+1	MAIN	180
		IF (CY.LE.0.) IER=IER+1	MAIN	181
195		IF (EN.LE.0.) IER=IER+1	MAIN	182
		PRINT 940, CY,CZ,EN	MAIN	183
	390	IF (DIAM.GT.0.) CALL DEP	MAIN	184
		IF (D5.GT.0.) GO TO 400	MAIN	185
		IF (D4.GT.0.) GO TO 400	MAIN	186
200		IF (D3.GT.0.) GO TO 400	MAIN	187
		IER=IER+1	MAIN	188
	400	PRINT 830,DIAM,D3,D4,D5	MAIN	189
		IF (R(1).LE.0.) IER=IER+1	MAIN	190
		IF (DOSTIM(1).LE.J.) IER=IER+1	MAIN	191
205		PRINT 950, NR,(R(I),I=1,NR)	MAIN	192
		PRINT 960, NUPT,(UPTIME(I),I=1,NUPT)	MAIN	193
		PRINT 970, NDT,(DOSTIM(I),I=1,NOT)	MAIN	194
		IF (IER.LT.1) GO TO 410	MAIN	195
		PRINT 1020, IER	MAIN	196
210		STOP	MAIN	197
	410	CONTINUE	MAIN	198
		OUB=1/UBAR	MAIN	199
		IGC=MET	MAIN	200
		GO TO (420,450,540,480,480,480,480,480,480,590), MET	MAIN	201
215	C	HANFORD MODEL	MAIN	202
	420	CONTINUE	MAIN	203
		DO 440 IU=1,NUPT	MAIN	204
		DO 430 IR=1,NR	MAIN	205
		TT=R(IR)*OUB	MAIN	206
220		TT2=TT*TT	MAIN	207
		EXPO=KSQD*TT2	MAIN	208
		EXX=EXP(-EXPO)	MAIN	209
		OME=1.-EXX	MAIN	210
		SZ2=BH*TT+OME*AH	MAIN	211
225		SIGZ(IR,IU)=SQRT(SZ2)	MAIN	212
		BA=CH+DH*S TUB(IU)	MAIN	213
		APH=.5*BA/(STUB(IU)**2)	MAIN	214
		SY2=BA*(TT-APH*(1.-EXP(-TT/APH)))	MAIN	215

		SIGY(IR,IU)=SQRT(SY2)	MAIN	215
230	430	CONTINUE	MAIN	217
	440	CONTINUE	MAIN	218
		GO TO 540	MAIN	219
	C	SUTTON MODEL	MAIN	220
	450	CONTINUE	MAIN	221
235		SQ2=1./SQRT(2.)	MAIN	222
		DO 470 IR=1,NR	MAIN	223
		SIGY(IR,1)=CY*SQ2*(R(IR)**(1.-.5*EN))	MAIN	224
		SIGZ(IR,1)=CZ*SQ2*(R(IR)**(1.-.5*EN))	MAIN	225
		DO 460 IU=2,NUPT	MAIN	226
240		SIGZ(IR,IU)=SIGZ(IR,1)	MAIN	227
		SIGY(IR,IU)=SIGY(IR,1)	MAIN	228
	460	CONTINUE	MAIN	229
	470	CONTINUE	MAIN	230
		GO TO 540	MAIN	231
245	C	PASQUILL CURVES FOR SIGMA Z	MAIN	232
	480	ITM3=MET-3	MAIN	233
		DO 490 IR=1,NR	MAIN	234
		X(IR)=100.*R(IR)	MAIN	235
	490	CONTINUE	MAIN	236
250		ERR=0.	MAIN	237
		CALL PASSIG (ITM3,NR,ERR)	MAIN	238
		IF (ERR) 510,510,500	MAIN	239
	500	WRITE (6,980) MET	MAIN	240
	510	CONTINUE	MAIN	241
255		DO 530 IR=1,NR	MAIN	242
		DO 520 IU=1,NUPT	MAIN	243
		SIGZ(IR,IU)=SZ(IR)/100.	MAIN	244
		SIGY(IR,IU)=SY(IR)/100.	MAIN	245
	520	CONTINUE	MAIN	245
260	530	CONTINUE	MAIN	247
	C	PRINT SIGMA Y AND SIGMA Z VALUES IF IPR GT 0	MAIN	248
	540	IF (IPR) 560,560,550	MAIN	249
	550	WRITE (6,990)	MAIN	250
		WRITE (6,1000) R	MAIN	251
265		WRITE (6,1010) UBAR,((SIGZ(IR,IU),IR=1,10),IU=1,NUPT)	MAIN	252
		WRITE (6,1070)	MAIN	253
		WRITE (6,1000) R	MAIN	254
		WRITE (6,1010) UBAR,((SIGY(IR,IU),IR=1,10),IU=1,NUPT)	MAIN	255
	560	CONTINUE	MAIN	256
270	C	CALCULATE EQQ	MAIN	257
		DO 580 IR=1,NR	MAIN	258
		DO 570 IU=1,NUPT	MAIN	259
		DENOM=SIGY(IR,IU)*SIGZ(IR,IU)*UBAR*3.1415927	MAIN	260
		EXPO=(-.5*(H/SIGZ(IR,IU))**2)	MAIN	261
275		EQQ(IR,IU)=EXP(EXPO)/DENOM	MAIN	262
	570	CONTINUE	MAIN	263
	580	CONTINUE	MAIN	264
		IF (IPU,GT.0) PUNCH 840, ((EQQ(I,J),I=1,10),J=1,NUPT)	MAIN	265
		GO TO 600	MAIN	266
280	590	READ (5,840) ((EQQ(I,J),I=1,10),J=1,NUPT)	MAIN	267
	600	CONTINUE	MAIN	268
	C	PRINT EQQ VALUES IF IQP.GT.0	MAIN	269
		IF (IQP.LE.0) GO TO 620	MAIN	270
		PRINT 820, IPAGE, JOY	MAIN	271
285		PRINT 1030, (R(I),I=1,NR)	MAIN	272

		IPAGE=IPAGE+1	MAIN	273
		DO 610 IU=1,NUPT	MAIN	274
		PRINT 1040, UPTIME(IU), (EQQ(IR,IU),IR=1,NR)	MAIN	275
	610	CONTINUE	MAIN	276
290	C	START DOSE CALCULATION	MAIN	277
	C	PERFORM DOSE CALCULATION	MAIN	278
	C	MAIN DO LOOP ON DISTANCES	MAIN	279
	620	DO 720 IR=1,NR	MAIN	280
		KR=IR	MAIN	281
295		TT=P(IR)/UBAR	MAIN	282
		IF (.NOT.FISPRD) GO TO 630	MAIN	283
		CALL CHAINT (2,TT,1)	MAIN	284
	630	IF (.NOT.ACTPRD) GO TO 650	MAIN	285
		DO 640 ISO=N1,NTOT	MAIN	286
300		IF (MASS(ISO,1).LT.1.E-30) GO TO 640	MAIN	287
		MASS(ISO,2)=MASS(ISO,1)*EXP(-LAM3DA(ISO)*TT)	MAIN	288
	640	CONTINUE	MAIN	289
	650	DO 660 ISO=1,NTOT	MAIN	290
	660	SMASS(ISO)=MASS(ISO,2)	MAIN	291
305		DO 710 IU=1,NUPT	MAIN	292
		BOU=BRATE(IU)*EQQ(IR,IU)/UPTIME(IU)	MAIN	293
		IF (.NOT.FISPRD) GO TO 680	MAIN	294
		DO 670 ISO=1,NFF	MAIN	295
	670	MASS(ISO,2)=SMASS(ISO)*BOU	MAIN	296
310	680	IF (.NOT.ACTPRD) GO TO 700	MAIN	297
		DO 690 ISO=N1,NTOT	MAIN	298
	690	MASS(ISO,2)=SMASS(ISO)*BOU	MAIN	299
	700	TLN=DTLN(IU)	MAIN	300
		TLN=DTLN(IU)	MAIN	301
315		KUPT=IU	MAIN	302
		CALL ORDDOSE	MAIN	303
	C	END OF CASE	MAIN	304
	710	CONTINUE	MAIN	305
	720	CONTINUE	MAIN	306
320		WRITE(1,725)	9/4/74	11
	725	FORMAT(" START OF NEXT CASE")	9/4/74	12
		DO 729 KO=1,IO	9/4/74	13
		DO 729 IO=1,NOT	9/4/74	14
		KOG=ORGANS(KO)	9/4/74	15
325		WRITE(1,728) TITLE2(1),IO,KOG,(SUMDOS(IR,IL,KO),IR=1,NR)	9/4/74	16
	728	FORMAT(A4,3X,I1,I2,10E7.2)	9/4/74	17
	729	CONTINUE	9/4/74	18
	C	RETURN TO START FOR NEXT CASE	MAIN	307
	C	CALCULATE WEIGHTED DOSE VALUES OVER TOTAL RELEASE PERIOD	MAIN	308
330		IF (NUPT.LT.2) GO TO 100	MAIN	309
		LFMT(2)=ISPACE(IO)	MAIN	310
		LFMT(7)=IFIELD(IO)	MAIN	311
		MFMT(2)=ISPACE(IO)	MAIN	312
		NFMT(2)=ISPACE(IO)	MAIN	313
335		JFMT(7)=IFIELD(IO)	MAIN	314
		JFMT(2)=ISPACE(IO)	MAIN	315
		DO 760 IU=2,NUPT	MAIN	316
		FAC=UPTIME(IU)/(UPTIME(IU)+UPTIME(IU-1))	MAIN	317
		DO 750 IO=1,NOT	MAIN	318
340		OU 740 KO=1,IO	MAIN	319
		DO 730 IR=1,NR	MAIN	320
		IRIU=IR + (IU-1)*10	MAIN	321

		IRIU=IR + (IU-2)*10	MAIN	322
		SUMDOS (IRIU,IO,KO)=SUMDOS (IRIU,IO,KO)*FAC+SUMDOS (IRIU,IO,KO)*(1.-	MAIN	323
345		1FAC)	MAIN	324
	730	CONTINUE	MAIN	325
	740	CONTINUE	MAIN	326
	750	CONTINUE	MAIN	327
	760	CONTINUE	MAIN	328
350		CALL UNIT (CTLM,CTLN,NOT,DOSTIM)	MAIN	329
		ILINE=40	MAIN	330
		DO 800 IU=1,NUPT	MAIN	331
		DO 790 IR=1,NR	MAIN	332
		IRIU=IR + (IU-1)*10	MAIN	333
355		IF (ILINE+NOT+3.LE.41) GO TO 770	MAIN	334
		ILINE=0	MAIN	335
		PRINT 1080,IPAGE,JOY,DTLM(IU),DTLN(IU)	MAIN	336
		WRITE (6,1090) ((ORGAN(I,J),I=1,2),J=1,IO)	MAIN	337
		IPAGE=IPAGE+1	MAIN	338
360	770	WRITE (6,1100) R(IR)	MAIN	339
		ILINE=ILINE+3	MAIN	340
		DO 780 IO=1,NOT	MAIN	341
		WRITE (6,1110) CTLM(IO),CTLN(IO),(SUMDOS (IRIU,IO,K),K=1,10)	MAIN	342
	780	CONTINUE	MAIN	343
365		ILINE=ILINE+NOT	MAIN	344
	790	CONTINUE	MAIN	345
	800	CONTINUE	MAIN	346
		GO TO 100	MAIN	347
	C	STOP ON END OF FILE	MAIN	348
370	310	STOP	MAIN	349
	C		MAIN	350
	C	ORGAN DOSE CALCULATION ONLY	MAIN	351
	815	IF (IER.GT.2) GO TO 819	MAIN	352
		PRINT 960,NUPT,(UPTIME(I),I=1,NUPT)	MAIN	353
375		PRINT 970,NOT,(DOSTIM(I),I=1,NOT)	MAIN	354
		KUPT=1	MAIN	355
		KR=1	MAIN	356
		DO 817 I=1,NUPT	MAIN	357
		DO 816 J=1,NOT	MAIN	358
380	816	MASS(J,2)=MASS(J,1)*1.E+6/UPTIME(I)	MAIN	359
		TLH=JTLH(I)	MAIN	360
		TLN=DTLN(I)	MAIN	361
		KUPT=I	MAIN	362
		KR=1	MAIN	363
385		CALL ORDOSE	MAIN	364
	817	CONTINUE	MAIN	365
		GO TO 100	MAIN	366
	819	IER=IER-1	MAIN	367
		PRINT 1020,IER	MAIN	368
390		STOP	MAIN	369
	C		MAIN	370
	820	FORMAT (1H1,"PAGE",I5,113X,A9//)	MAIN	371
	830	FORMAT (1H,"DIAM =",F3.3," MICROMETERS"/1H,"D3 =",F9.3/1H,"D4 =",	MAIN	372
		1",F9.3/1H,"D5 =",F9.3)	MAIN	373
395	840	FORMAT (10E7.3)	MAIN	374
	850	FORMAT (I3)	MAIN	375
	855	FORMAT (1H1,"PAGE",I5,45X,"***** INPUT DATA *****",44X,A9///)	MAIN	376
	856	FORMAT (1H0,7A10)	MAIN	19
	860	FORMAT (1H0,53X,"NUCLIDE RELEASE IN CURIES"//)	MAIN	377

400	870	FORMAT(1H ,2X,6(2X,A3,I3,I2,1PE11.4))	MAIN	378
	880	FORMAT (4(A3,I3,I2,E12.0))	MAIN	379
	890	FORMAT ("UNKNOWN ISOTOPE - "A3,I3)	MAIN	380
	900	FORMAT (" INVALID VALUE FOR NEXT - "16,)	MAIN	381
	910	FORMAT (" NO ORGANS SPECIFIED - ORGANS",10I2)	MAIN	382
405	920	FORMAT (1H0///1HG,"UBAR =",F6.1/1H ,"DRATE =",5F9.0/1H ,"MET =",I3)	MAIN	383
	930	FORMAT (" AH="F7.1/" BH="F8.5/" CH="F7.1/" DH="F7.1/" KSQD="F9.6/	MAIN	384
		1" STUB(1)="F7.3)	MAIN	385
	940	FORMAT (" CY="F6.3/" CZ="F6.3/" EN="F6.3)	MAIN	386
	950	FORMAT (" "I3," DISTANCES, METERS"10F10.0)	MAIN	387
410	960	FORMAT (1H ,I3," UPTAKE TIMES, SECONDS ",1P5E10.3)	MAIN	388
	970	FORMAT (1H ,I3," DOSE TIMES, SECONDS ",1P10E10.3)	MAIN	389
	980	FORMAT (57HGLIMITS OF STORED SIGZ DATA EXCEEDED FOR METEOROLOGY TY	MAIN	390
		1PE,I3)	MAIN	391
	990	FORMAT (//60X,"SIGMA Z VALUES")	MAIN	392
415	1000	FORMAT (13H0AVERAGE WIND,3CX,34HDISTANCE FROM RELEASE POINT METERS	MAIN	393
		1/13H SPEED, M/SEC,10F10.0/)	MAIN	394
	1010	FORMAT (1H ,F10.1,2X,10F10.0/(1H ,12X,10F10.0))	MAIN	395
	1020	FORMAT (" IER = "I3," ERRORS IN INPUT DATA - STOP")	MAIN	396
	1030	FORMAT (1HG,51X,"E/Q VALUES"/1H0, "UPTAK	MAIN	397
420		1 TIMES / DISTANCES"/1H ,2X,"(SECONDS)",EX,"(METERS)",10F10.0)	MAIN	398
	1040	FORMAT (1HG,1PE10.3,15X,10E10.3/(1H ,1PE10.3,15X,10E10.3))	MAIN	399
	1050	FORMAT (8A1C)	MAIN	400
	1060	FORMAT (" IMPROPER SOLUBILITY INDEX SPECIFIED FOR ",A3,I3,I2)	MAIN	401
	1070	FORMAT (//60X,"SIGMA Y VALUES")	MAIN	402
425	1080	FORMAT (1H1,"PAGE",IE,42X,"DOSE COMMITMENT FROM UPTAKE",4-X,A9/	MAIN	403
		11H ,40X,"AVERAGED OVER A TOTAL RELEASE PERIOD OF ",F6.1,1X,A6/1H0,	MAIN	404
		249X,"TASK GROUP ON LUNG DYNAMICS MODEL"/1H ,56X,"ORGAN DOSE IN REM	MAIN	405
		3"/)	MAIN	406
	1090	FORMAT ("DDISTANCE/"33X"ORGANS CONSIDERED ARE"/" DOSE TIME "10(MAIN	407
430		1A6,A5)	MAIN	408
	1100	FORMAT ("0"F8.0," METERS"/)	MAIN	409
	1110	FORMAT (" "F5.1,1XA5,1P10E11.2)	MAIN	410
		END	MAIN	411

		SUBROUTINE LIBB	LIBB	2
		LIBB READS AN ORGAN DATA LIBRARY.	LIBB	3
		COMMON NFP,NAC,NTOT,NDFNUC(96),SKIP(96),MASS(600,2),LAMBDA(620),	COMMON	2
		1 CHAINS,OKFRCT(450,3),PEC(600),ML(600),ICL(600)	COMMON	3
5		COMMON MA(600),LAMBIO(1200),FMAT(1200),FAIR(1200),F2P(1200)	COMMON	4
		COMMON IORG(1200),ENG(1200,3),F1(600),IBEG(600),IEND(600)	COMMON	5
		COMMON DOSTIM(13),D3,D4,D5,UPT,ML(5),ISM(600)	COMMON	6
		COMMON BRATE(5),R(10),X(10),SY(13),SZ(10),DIAM	COMMON	7
		COMMON SUMDOS(50,10,10),TITLE2(8),NOT,KORG(20),TLM,TLN	COMMON	8
10		COMMON IPAGE,ORGAN(2,10)	COMMON	9
		COMMON/NAME/ORGNAME(2,25)	COMMON	10
		REAL LAMBIO,LAMBDA,MASS	COMMON	11
		INTEGER CHAINS,SKIP	COMMON	12
		LOGICAL FISPRD,ACTPRD	COMMON	13
15		COMMON ACIPRO,FISPRD,KR,KUPT,DOY	COMMON	14
		COMMON LIBT(7)	9/9/74	1
		LOGICAL SKIPO	LIBB	5
		C READ BIOLOGICAL DATA LIBRARY	LIBB	6
	1	FORMAT(I4,7A10)	9/4/74	1
20	2	FORMAT(2I3,I1,A2,I2,6E8,3,2E6,2)	LIBB	8
	3	FORMAT(" LIBRARY DATA SEQUENCE ERROR "// "2I3,I1,A2,I2,6E10,4)	LIBB	9
		SKIPO=.FALSE.	LIBB	10
		DO 100 I=1,600	LIBB	11
	100	IBEG(I)=0	LIBB	12
25		KRN=0	LIBB	13
		KET=0	LIBB	14
		KIST=0	LIBB	15
		KMT=0	LIBB	16
		IOT=0	LIBB	17
30		KJ=0	LIBB	18
		READ(28,1) NCOS,(LIBT(K),K=1,7)	9/9/74	2
		DO 300 I=1,NCOS	LIBB	20
		READ(28,2) KE,KIS,KM,ELT,IO,TR,T3,FM,FA,FP2,EFF1,EFF2,EFF3	LIBB	21
		IF(KE-KET) 150,110,160	LIBB	22
35	110	IF(KIS-KIST) 150,120,160	LIBB	23
	120	IF(KM-KMT) 150,130,160	LIBB	24
	130	IF(IO-IOT) 150,150,210	LIBB	25
		C LIBRARY DATA SEQUENCE ERROR	LIBB	26
40	150	PRINT 3, KE,KIS,KM,ELT,IO,TR,T3,FM,FA,FP2,EFF1,EFF2,EFF3	LIBB	27
		STOP	LIBB	28
	160	IOT=1	LIBB	29
		KET=KE	LIBB	30
		KIST=KIS	LIBB	31
		KMT=KM	LIBB	32
45		IF(IO-1) 150,170,150	LIBB	33
	170	IF(KRN.GT.C) IEND(KJ)=KRN	LIBB	34
		SKIPO=.FALSE.	LIBB	35
		DO 190 J=1,NTOT	LIBB	36
		IF(KE.NE.MA(J).OR.KIS.NE.ML(J)) GO TO 190	LIBB	37
50		IF(KM) 175,175,180	LIBB	38
	175	IF(KM+2.EQ.ISM(J)) GO TO 200	LIBB	39
		GO TO 190	LIBB	40
	180	IF(KM.EQ.ISM(J)) GO TO 200	LIBB	41
	190	CONTINUE	LIBB	42
55		C ISOTOPE NOT IN R130 LIBRARY	LIBB	43
		SKIPO=.TRUE.	LIBB	44
		GO TO J00	LIBB	45

SUBROUTINE LIB8

74/74

OPT=C

TRACE

FTN 4.2+P380

12/10/74 1..38.40.

	C NEW ISOTOPE	LIB8	46
	200 KJ=J	LIB8	47
60	IBEG(KJ)=KRN+1	LIB8	48
	F1(KJ)=FM	LIB8	49
	210 IF(SKIP0) GO TO 300	LIB8	50
	IOT=IO	LIB8	51
	KRN=KRN+1	LIB8	52
65	IF(TB.LE.0.) GO TO 220	LIB8	53
	LAMBIO(KRN)=.69314/(96400.*TB)	LIB8	54
	GO TO 230	LIB8	55
	220 LAMBIO(KRN)=0.	LIB8	56
70	230 FMAT(KRN)=FM	LIB8	57
	FAIR(KRN)=FA	LIB8	58
	F2P(KRN)=FP2	LIB8	59
	IORG(KRN)= IO	LIB8	60
	ENG(KRN,1)=EFF1	LIB8	61
	ENG(KRN,2)=EFF2	LIB8	62
75	ENG(KRN,3)=EFF3	LIB8	63
	IF(EFF2.LE.0.) ENG(KRN,2)=EFF1	LIB8	64
	IF(EFF3.LE.0.) ENG(KRN,3)=EFF1	LIB8	65
	300 CONTINUE	LIB8	66
	IEND(KJ)=KRN	LIB8	67
80	RETURN	LIB8	68
	END	LIB8	69

		SUBROUTINE LIBK	LIBK	2
	C	LIBK READS AN ISOTOPE DATA LIBRARY	LIBK	3
		COMMON NFP,NAC,NTOT,NOFNUC(96),SKIP(96),MASS(600,2),LAMBDA(600),	COMN	2
		1 CHAINS,DKFRCT(450,3),REC(600),ML(600),ICL(600)	COMN	3
5		COMMON MA(600),LAMBIO(1200),FMAT(1200),FAIR(1200),F2P(1200)	COMN	4
		COMMON IORG(1200),ENG(1200,3),F1(600),ISEG(600),IEND(600)	COMN	5
		COMMON DOSTIM(10),D3,04,05,UPTIM(5),ISM(600)	COMN	6
		COMMON BRATE(5),R(10),X(10),SY(10),SZ(10),DIAM	COMN	7
		COMMON SUMDOS(50,10,10),TITLE2(8),NDT,KORG(20),TLH,TLN	COMN	8
10		COMMON IPAGE,ORGAN(2,10)	COMN	9
		COMMON/NAM/ORGNAM(2,25)	COMN	10
		REAL LAMBIO,LAMBDA,MASS	COMN	11
		INTEGER CHAINS,SKIP	COMN	12
		LOGICAL FISPRO,ACTPRO	COMN	13
15		COMMON ACTPRO,FISPRO,KR,KUPT,DOY	COMN	14
		DIMENSION FRACT(450)	LIBK	5
		LOGICAL ERROR	LIBK	6
	C		LIBK	7
	C	READ RIBD DATA LIBRARY	LIBK	8
20		READ(29,1) NCD	LIBK	9
	1	FORMAT(I3)	LIBK	10
		READ(29,2) (ML(J),MA(J),LAMBDA(J),FRACT(J),REC(J),ISM(J),J=1,NCD)	LIBK	11
	2	FORMAT(I3,I2,E9.3,32X,F5.3,17X,A3,I1)	LIBK	12
		READ(29,1) N2	LIBK	13
25		N1=NCD+1	LIBK	14
		NAC=N2	LIBK	15
		N2=N2+NCD	LIBK	16
		NFP=NCD	LIBK	17
		NTOT=N2	LIBK	18
30		READ(29,102) (ML(J),MA(J),LAMBDA(J),REC(J),ISM(J),J=N1,N2)	LIBK	19
	102	FORMAT(I3,I2,E9.3,54X,A3,I1)	LIBK	20
		DO 202 L=N1,N2	LIBK	21
		IF(LAMBDA(L).LT.1.E-30) GO TO 202	LIBK	22
		LAMBDA(L)=.69314/86400./LAMBDA(L)	LIBK	23
35	202	CONTINUE	LIBK	24
		ERROR=.FALSE.	LIBK	25
		MP=0	LIBK	26
		CHAINS=0	LIBK	27
		DO 3 I=1,96	LIBK	28
40	3	NOFNUC(I)=0	LIBK	29
	C	CALCULATE LAMBDA IN SEC-1	LIBK	30
		DO 10 J=1,NCD	LIBK	31
		IF(LAMBDA(J).LT.1.E-30) GO TO 40	LIBK	32
		LAMBDA(J)=.69314/LAMBDA(J)/86400.	LIBK	33
45	40	IF(MP-ML(J)) 4,5,6	LIBK	34
	4	MP=ML(J)	LIBK	35
	C	NEW MASS CHAIN	LIBK	36
		CHAINS=CHAINS+1	LIBK	37
50	C	COUNT NUMBER OF NUCLIDES IN THIS CHAIN	LIBK	38
	5	NOFNUC(CHAINS)=NOFNUC(CHAINS)+1	LIBK	39
		GO TO 8	LIBK	40
	6	ERROR=.TRUE.	LIBK	41
		PRINT 7,J	LIBK	42
	7	FORMAT("3LIBRARY ENTRY"4," IS OUT OF PLACE")	LIBK	43
55	8	DO 9 L=1,3	LIBK	44
	9	DKFRCT(J,L)=0.0	LIBK	45
	10	CONTINUE	LIBK	46

		IF(IECR) STOP	LIBK	47
	C	ISM=1 MEANS ISOMERIC STATE,=2 MEANS GROUND STATE	LIBK	48
60		ISOTOP=0	LIBK	49
		DO 15 K=1,CHAINS	LIBK	50
		LIN=NOFNUC(K)	LIBK	51
		J=0	LIBK	52
	C	DETERMINE FRACTIONAL DECAY YIELDS	LIBK	53
65		DO 14 I=1,LIM	LIBK	54
		J=J+1	LIBK	55
		ISOTOP=ISOTOP+1	LIBK	56
		IF(J.EQ.1) GO TO 13	LIBK	57
		IF(ISM(ISOTOP).EQ.2) GO TO 11	LIBK	58
70		DKFRCT(ISOTOP,1)=FRACT(ISOTOP-1)	LIBK	59
		GO TO 13	LIBK	60
	11	DKFRCT(ISOTOP,1)=1.-FRACT(ISOTOP-1)	LIBK	61
		IF(J.EQ.2) GO TO 13	LIBK	62
		IF(ISM(ISOTOP-1).EQ.1) GO TO 12	LIBK	63
75		IF(ISM(ISOTOP-2).EQ.2) GO TO 13	LIBK	64
		DKFRCT(ISOTOP,2)=FRACT(ISOTOP-2)	LIBK	65
		GO TO 13	LIBK	66
	12	DKFRCT(ISOTOP,2)=1.-FRACT(ISOTOP-2)	LIBK	67
		IF(J.EQ.3.OR.ISM(ISOTOP-3).EQ.2) GO TO 13	LIBK	68
80		DKFRCT(ISOTOP,3)=FRACT(ISOTOP-3)	LIBK	69
	13	IF(LAMBDA(ISOTOP).LT.1.E-30) J=0	LIBK	70
	14	CONTINUE	LIBK	71
	15	CONTINUE	LIBK	72
	C	RETURN	LIBK	73
85		END	LIBK	74
			LIBK	75

		SUBROUTINE DEP	DEP	2
	C	DEP CALCULATES THE RESPIRATORY TRACT DEPOSITION FACTORS FROM THE	DEP	3
	C	ACTIVITY, MEDIAN ATMOSPHERIC DIAMETER OF AN AEROSOL DISTRIBUTION	DEP	4
		COMMON NFP, NAC, NTOT, NUFNUC(96), SKIP(96), MASS(600,2), LAMBDA(500),	COMN	2
5		1 CHAINS, DKFRCT(450,3), REC(600), NL(600), ICL(600)	COMN	3
		COMMON MA(500), LAMBIG(1200), FNAT(1200), FAIR(1200), F2P(1200)	COMN	4
		COMMON IORG(1200), ENG(1200,3), F1(600), IBEG(600), IEND(600)	COMN	5
		COMMON DOSTIM(10), D3, D4, D5, UPTIME(5), ISM(600)	COMN	6
		COMMON BKATE(5), R(10), X(10), SY(10), SZ(10), DIAM	COMN	7
10		COMMON SUMDOS(50,10,10), TITLE2(8), NOT, KOR6(20), TLH, TLN	COMN	8
		COMMON IPAGE, ORGAM(2,10)	COMN	9
		COMMON/NAM/ORGNA(12,25)	COMN	10
		REAL LAMBIG, LAMBDA, MASS	COMN	11
		INTEGER CHAINS, SKIP	COMN	12
15		LOGICAL FISPRD, ACTPRD	COMN	13
		COMMON ACTFRD, FISPRD, KR, KUPT, DOY	COMN	14
		EXTERNAL ERFXP	DEP	6
		DELTA=.000201	DEP	7
		DP=.1	DEP	8
20		SP=24.	DEP	9
		DPL=ALOG10(DP)	DEP	10
		SPL=ALOG10(SP)	DEP	11
		XL=ALOG10(DIAM)	DEP	12
		XP=(XL-DPL)/SPL	DEP	13
25		FF=SIMPS(-5.0, XP, DELTA, ERFXP)	DEP	14
		FAC=SQRT(2.*3.1415927)	DEP	15
		D3=1.-FP/FAC	DEP	16
		DP=2.	DEP	17
		SP=3.45	DEP	18
30		DPL=ALOG10(DP)	DEP	19
		SPL=ALOG10(SP)	DEP	20
		XP=(XL-DPL)/SPL	DEP	21
		FP=SIMPS(-5.0, XP, DELTA, ERFXP)	DEP	22
		D3=FP/FAC	DEP	23
35		D4=.08	DEP	24
		RETURN	DEP	25
		END	DEP	26

FUNCTION ERFXP 74774 OPT=0 TRACE

FTN 4.2+P380

12/10/74 14.29.23.

C
5
FUNCTION ERFXP(ARG)
ERFXP IS THE ERROR FUNCTION
EX=-.5*(ARG**2)
ERFXP=EXP(EX)
RETURN
END

ERFXP 2
E<FXP 3
E<FXP 4
ERFXP 5
ERFXP 6
ERFXP 7

	FUNCTION SIMPS(A,B,DELTA,FC0000)	SIMPS	2
C	A AND B ARE MIN AND MAX OF THE DEFINITE INTEGRAL	SIMPS	3
C	DELTA IS THE PERMISSIBLE DIFFERENCE BETWEEN TWO SUCCESSIVE SUMS.	SIMPS	4
	N=1	SIMPS	5
5	H=(B-A)/2.	SIMPS	6
	FJAY=H*(FC0000(A)+FC0000(B))	SIMPS	7
C	A MAXIMUM OF 2049 POINTS WILL BE USED WHEN L=11	SIMPS	8
	DO 35 L=1,11	SIMPS	9
10	S=0.0	SIMPS	10
	DO 20 K=1,N	SIMPS	11
10	S=S+FC0000(A+(2.0*FLOAT(K)-1.0)*H)	SIMPS	12
	FI=FJAY+4.0*H*S	SIMPS	13
	IF(L-3)26,26,23	SIMPS	14
	23 CONTINUE	SIMPS	15
15	IF(DELTA-ABS((FI-BAR)/FI))30,40,+0	SIMPS	16
	26 CONTINUE	SIMPS	17
	30 BAR=FI	SIMPS	18
	FJAY=(FI+FJAY)/4.0	SIMPS	19
	N=2*N	SIMPS	20
20	H=H/2.0	SIMPS	21
	35 CONTINUE	SIMPS	22
	40 SIMPS =FI/3.0	SIMPS	23
	RETURN	SIMPS	24
	END	SIMPS	25

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SUBROUTINE PASSIG(TYPE,NUM,ERR)
C PASSIG INTERPOLATES STORED PASQUILL CURVES FOR SY AND SZ FOR TYPE TYP
C AND NUM X POSITIONS
COMMON NFP,NAC,NTOT,NOFNUC(96),SKIP(96),MASS(600,2),LAMBDA(500),
1CHAINS,OKFRCT(450,3),REC(500),ML(600),ICL(600)
COMMON MA(600),LAMBDA(1200),FMAT(1200),FAIR(1200),F2P(1200)
COMMON IORG(1200),ENG(1200,3),F1(600),IBEG(600),IEND(600)
COMMON DOSTIM(10),D3,D4,D5,UPTIME(5),ISM(600)
COMMON GRATE(5),R(10),X(10),SY(10),SZ(10),DIAM
COMMON SUMLOS(50,10,10),TITLE2(8),NOT,KORG(20),TLH,TLN
COMMON IPAGE,ORGAN(2,10)
COMMON/NAM/ORGNA(2,25)
REAL LAMBIC,LAMBDA,MASS
INTEGER CHAINS,SKIP
LOGICAL FISPRD,ACTPRD
COMMON ACTPRD,FISPRD,KR,KUPT,DOY
INTEGER TYPE
DIMENSION SIGY(6,20),SIGZ(6,20),DIST(20)
C DISTANCES TO WHICH SY AND SZ DATA CORRESPOND.
DATA DIST/C.0,1.E4,1.5E4,2.5E4,3.5E4,5.E4,7.E4,1.E5,1.5E5,2.5E5,
13.5E5,5.E5,7.E5,1.E6,1.5E6,2.5E6,3.5E6,5.E6,7.E6,1.E7/
C SIGMA Y DATA - 1 FOR TYPE A, 2 FOR TYPE B, ETC.
DATA (SIGY(1,I),I=1,20)/20.,2.75E3,3.8E3,6.1E3,8.3E3,1.15E4,1.54E4,
1.2.14E4,3.05E4,4.7E4,6.3E4,8.7E4,1.14E5,1.36E5,2.2E5,3.3E5,4.4E5,
25.95E5,9.E5,1.15E6/
DATA (SIGY(2,I),I=1,20)/90.,1.95E3,2.75E3,4.4E3,6.E3,8.3E3,1.12E4,
11.56E4,2.25E4,3.6E4,4.9E4,6.6E4,8.8E4,1.18E5,1.68E5,2.6E5,3.45E5,4
2.6E5,6.1E5,8.2E5/
DATA (SIGY(3,I),I=1,20)/90.,1.26E3,1.8E3,2.9E3,4.E3,5.5E3,7.5E3,1.
104E4,1.51E4,2.44E4,3.3E4,4.5E4,6.1E4,8.4E4,1.19E5,1.87E5,2.5E5,3.3
25E5,4.5E5,6.2E5/
DATA (SIGY(4,I),I=1,20)/90.,8.2E2,1.18E3,1.9E3,2.6E3,3.6E3,5.9E3,6
1.9E3,1.E4,1.6E4,2.17E4,3.E4,4.05E4,5.5E4,7.9E4,1.21E5,1.62E5,2.2E5
2.3.E5,4.1E5/
DATA (SIGY(5,I),I=1,20)/90.,6.1E2,8.8E2,1.4E3,1.9E3,2.6E3,3.6E3,5.
105E3,7.3E3,1.18E4,1.6E4,2.22E4,3.E4,4.2E4,5.9E4,9.2E4,1.21E5,1.63E
25,2.25E5,3.E5/
DATA (SIGY(6,I),I=1,20)/90.,4.2E2,5.8E2,9.4E2,1.19E3,1.77E3,2.4E3,
13.4E3,4.9E3,7.8E3,1.07E4,1.48E4,2.E4,2.72E4,3.95E4,6.2E4,8.3E4,1.1
22E5,1.5E5,2.E5/
C SIGMA Z DATA - 1 FOR TYPE A, 2 FOR TYPE B, ETC.
DATA (SIGZ(1,I),I=1,20)/100.,1.4E3,2.2E3,3.8E3,6.E3,1.08E4,2.17E4,
14.6E4,1.68E5,3.15E5,6.0E5,9*-1.0/
DATA (SIGZ(2,I),I=1,20)/100.,1.08E3,1.56E3,2.55E3,3.5E3,5.1E3,7.5E
13,1.1E4,1.7E4,3.E4,4.3E4,6.4E4,9.2E4,1.35E5,2.1E5,3.75E5,6.4E5,-1.
2,-1.,-1./
DATA (SIGZ(3,I),I=1,20)/100.,7.3E2,1.06E3,1.7E3,2.3E3,3.2E3,4.4E3,
16.1E3,8.9E3,1.42E4,1.95E4,2.65E4,3.65E4,5.E4,7.4E4,1.15E5,1.6E5,2.
22E5,2.9E5,4.E5/
DATA (SIGZ(4,I),I=1,20)/100.,4.6E2,6.6E2,1.02E3,1.37E3,1.8E3,2.4E
13,3.15E3,4.2E3,5.8E3,7.1E3,8.8E3,1.1E4,1.37E4,1.72E4,2.3E4,2.7E4,
23.3E4,3.9E4,4.6E4/
DATA (SIGZ(5,I),I=1,20)/100.,3.5E2,5.E2,7.6E2,9.9E2,1.3E3,1.65E3,2
1.15E3,2.8E3,3.8E3,4.7E3,5.5E3,6.6E3,7.9E3,9.6E3,1.21E4,1.39E4,1.54
2E4,1.7E4,1.85E4/
DATA (SIGZ(6,I),I=1,20)/100.,2.3E2,3.2E2,4.9E2,6.3E2,9.3E2,1.1E3,1
1.4E3,1.8E3,2.4E3,2.8E3,3.4E3,4.E3,4.65E3,5.4E3,6.4E3,7.1E3,7.8E3,

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		28.6E3,9.3E3/ IDATA=1	PASSIG 47
60	C	CALCULATE SY AND SZ FOR EACH X POSITION DESIRED	PASSIG 48
		DO 5 I=1,NUM	PASSIG 49
		1 IF(IDATA.GT.20) GO TO 7	PASSIG 50
		IF(SIGY(TYPE,IDATA)) 7,7,101	PASSIG 51
65	101	IF(SIGZ(TYPE,IDATA)) 7,7,201	PASSIG 52
		201 IF(X(I)-DIST(IDATA)) 4,3,2	PASSIG 53
		2 IDATA=IDATA+1	PASSIG 54
		GO TO 1	PASSIG 55
		3 SY(I)=SIGY(TYPE,IDATA)	PASSIG 56
70		SZ(I)=SIGZ(TYPE,IDATA)	PASSIG 57
		GO TO 5	PASSIG 58
		4 IF(IDATA.EQ.1) GO TO 3	PASSIG 59
		SY1=SIGY(TYPE,IDATA-1)	PASSIG 60
		SY2=SIGY(TYPE,IDATA)	PASSIG 61
75		SZ1=SIGZ(TYPE,IDATA-1)	PASSIG 62
		SZ2=SIGZ(TYPE,IDATA)	PASSIG 63
		D1=DIST(IDATA-1)	PASSIG 64
		D2=DIST(IDATA)	PASSIG 65
		SY(I)=SY1+(X(I)-D1)*(SY2-SY1)/(D2-D1)	PASSIG 66
80		SZ(I)=SZ1+(X(I)-D1)*(SZ2-SZ1)/(D2-D1)	PASSIG 67
		5 CONTINUE	PASSIG 68
		RETURN	PASSIG 69
		7 ERR=1.	PASSIG 70
		RETURN	PASSIG 71
85		END	PASSIG 72
			PASSIG 73
			PASSIG 74

		SUBROUTINE CHAINT(LOC,THETA,POSIT)	CHAINT	2
	C	SUBROUTINE CHAIN2 DECAYS NUCLIDES IN MASS ARRAY LOC FOR A TIME THETA	CHAINT	3
	C	STORES RESULTS IN MASS ARRAY POSIT	CHAINT	4
5	C	DIMENSION EXPO(8)	CHAINT	5
		COMMON NFP,NAC,NTOT,NOFNUC(96),SKIP(96),MASS(600,2),LAMBDA(600),	COMN	2
	1	CHAINS,DKFRCT(400,3),REC(600),ML(800),ICL(600)	COMN	3
		COMMON MA(600),LAMBDA(1200),FWAT(1200),FAIR(1200),F2P(1200)	COMN	4
		COMMON IORG(1200),ENG(1200,3),F1(600),I3EG(600),IEND(600)	COMN	5
10		COMMON DOSTIM(10),D3,D4,D5,UPTIME(5),ISY(600)	COMN	6
		COMMON BRATE(5),R(10),X(10),SY(10),SZ(10),DIAM	COMN	7
		COMMON SUMDOS(50,10,10),TITLE2(8),NOT,KORG(20),TLH,TLN	COMN	8
		COMMON IPAGE,ORGAN(2,10)	COMN	9
		COMMON/NAH/ORGNAH(2,25)	COMN	10
15		REAL LAMBDA,LAMBDA,MASS	COMN	11
		INTEGER CHAINS,SKIP	COMN	12
		LOGICAL FISPRD,ACTPRD	COMN	13
		COMMON ACTPRD,FISPRD,KR,KUPT,DJY	COMN	14
		INTEGER POSIT	CHAINT	8
20		ISOTOP=0	CHAINT	9
	C	DO LOOP ON NUMBER OF CHAINS	CHAINT	10
		DO 11 K=1,CHAINS	CHAINT	11
		LIM=NOFNUC(K)	CHAINT	12
		IF(SKIP(K).NE.0) GO TO 10	CHAINT	13
25		J=0	CHAINT	14
	C	DO LOOP ON ISOTOPES IN THE CHAIN	CHAINT	15
		DO 9 I=1,LIM	CHAINT	16
		ISOTOP=ISOTOP+1	CHAINT	17
		J=J+1	CHAINT	18
30		IF(LAMBDA(ISOTOP).LT.1.E-30) GO TO 8	CHAINT	19
		EXPO(J)=EXP(-LAMBDA(ISOTOP)*THETA)	CHAINT	20
	C	TRANSFER ON CHAIN MEMBER J	CHAINT	21
		GO TO (1,2,3,4,5,6,7),J	CHAINT	22
35	C	FIRST CHAIN MEMBER	CHAINT	23
		1 A1=MASS(ISOTOP,POSIT)	CHAINT	24
		MASS(ISOTOP,LOC)=A1*EXPO(1)	CHAINT	25
		GO TO 9	CHAINT	26
	C	SECOND CHAIN MEMBER	CHAINT	27
40		2 A2=DKFRCT(ISOTOP,1)*LAMBDA(ISOTOP-1)*A1/(LAMBDA(ISOTOP)-LAMBDA(ISO	CHAINT	28
		TOP-1))	CHAINT	29
		A3=MASS(ISOTOP,POSIT)-A2	CHAINT	30
		MASS(ISOTOP,LOC)=A3*EXPO(2)+A2*EXPO(1)	CHAINT	31
		GO TO 9	CHAINT	32
	C	THIRD CHAIN MEMBER	CHAINT	33
45		3 A4=DKFRCT(ISOTOP,1)*LAMBDA(ISOTOP-1)	CHAINT	34
		A4=(A4+DKFRCT(ISOTOP,2)*LAMBDA(ISOTOP-2)*A1)/(LAMBDA(ISOTOP)-	CHAINT	35
		LAMBDA(ISOTOP-2))	CHAINT	36
		A5=A4*A3/(LAMBDA(ISOTOP)-LAMBDA(ISOTOP-1))	CHAINT	37
		A6=MASS(ISOTOP,POSIT)-A4-A5	CHAINT	38
50		MASS(ISOTOP,LOC)=A6*EXPO(3)+A5*EXPO(2)+A4*EXPO(1)	CHAINT	39
		GO TO 9	CHAINT	40
	C	FOURTH CHAIN MEMBER	CHAINT	41
		4 A4=DKFRCT(ISOTOP,1)*LAMBDA(ISOTOP-1)	CHAINT	42
		BAKER=DKFRCT(ISOTOP,2)*LAMBDA(ISOTOP-2)	CHAINT	43
55		A7=(A4+BAKER*A2+DKFRCT(ISOTOP,3)*LAMBDA(ISOTOP-3)*A1)/(LAMBDA	CHAINT	44
		(ISOTOP)-LAMBDA(ISOTOP-3))	CHAINT	45
		A8=(A4+A5+BAKER*A3)/(LAMBDA(ISOTOP)-LAMBDA(ISOTOP-2))	CHAINT	46

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A9=ABLE*A6/(LAMBDA(ISOTOP)-LAMBDA(ISOTOP-1))          CHAINT 47
A10=MASS(ISOTOP,POSIT)-A9-A8-A7                       CHAINT 48
60 MASS(ISOTOP,LOC)=A10*EXPO(1)+A9*EXPO(3)+A8*EXPO(2)+A7*EXPO(1) CHAINT 49
GO TO 9                                                CHAINT 50
C FIFTH CHAIN MEMBER
5 ABLE=DKFRCT(ISOTOP,1)*LAMBDA(ISOTOP-1)              CHAINT 52
BAKER=DKFRCT(ISOTOP,2)*LAMBDA(ISOTOP-2)              CHAINT 53
65 CHARLY=DKFRCT(ISOTOP,3)*LAMBDA(ISOTOP-3)           CHAINT 54
A11=(ABLE*A7+BAKER*A4+CHARLY*A2)/(LAMBDA(ISOTOP)-LAMBDA(ISOTOP-4)) CHAINT 55
1) A12=(ABLE*A8+BAKER*A5+CHARLY*A3)/(LAMBDA(ISOTOP)-LAMBDA(ISOTOP-3)) CHAINT 56
1) A13=(ABLE*A9+BAKER*A6)/(LAMBDA(ISOTOP)-LAMBDA(ISOTOP-2)) CHAINT 58
70 A14=ABLE*A10/(LAMBDA(ISOTOP)-LAMBDA(ISOTOP-1)) CHAINT 60
A15=MASS(ISOTOP,POSIT)-A14-A13-A12-A11 CHAINT 61
MASS(ISOTOP,LOC)=A15*EXPO(5)+A14*EXPO(4)+A13*EXPO(3)+A12*EXPO(2)+ CHAINT 62
A11*EXPO(1) CHAINT 63
75 GO TO 9 CHAINT 64
C SIXTH CHAIN MEMBER
6 ABLE=DKFRCT(ISOTOP,1)*LAMBDA(ISOTOP-1) CHAINT 65
BAKER=DKFRCT(ISOTOP,2)*LAMBDA(ISOTOP-2) CHAINT 66
80 CHARLY=DKFRCT(ISOTOP,3)*LAMBDA(ISOTOP-3) CHAINT 67
A16=(ABLE*A11+BAKER*A7+CHARLY*A4)/(LAMBDA(ISOTOP)-LAMBDA(ISOTOP-5)) CHAINT 68
1) A17=(ABLE*A12+BAKER*A8+CHARLY*A5)/(LAMBDA(ISOTOP)-LAMBDA(ISOTOP-4)) CHAINT 69
1) A18=(ABLE*A13+BAKER*A9+CHARLY*A6)/(LAMBDA(ISOTOP)-LAMBDA(ISOTOP-3)) CHAINT 70
85 A19=(ABLE*A14+BAKER*A10)/(LAMBDA(ISOTOP)-LAMBDA(ISOTOP-2)) CHAINT 71
A20=ABLE*A15/(LAMBDA(ISOTOP)-LAMBDA(ISOTOP-1)) CHAINT 72
A21=MASS(ISOTOP,POSIT)-A20-A19-A18-A17-A16 CHAINT 73
MASS(ISOTOP,LOC)=A21*EXPO(6)+A20*EXPO(5)+A19*EXPO(4)+A18*EXPO(3) CHAINT 74
90 1+A17*EXPO(2)+A16*EXPO(1) CHAINT 75
GO TO 9 CHAINT 76
C SEVENTH CHAIN MEMBER
7 ABLE=DKFRCT(ISOTOP,1)*LAMBDA(ISOTOP-1) CHAINT 77
BAKER=DKFRCT(ISOTOP,2)*LAMBDA(ISOTOP-2) CHAINT 78
95 CHARLY=DKFRCT(ISOTOP,3)*LAMBDA(ISOTOP-3) CHAINT 79
A22=(ABLE*A16+BAKER*A11+CHARLY*A7)/(LAMBDA(ISOTOP)-LAMBDA(ISOTOP-6)) CHAINT 80
1) A23=(ABLE*A17+BAKER*A12+CHARLY*A8)/(LAMBDA(ISOTOP)-LAMBDA(ISOTOP-5)) CHAINT 81
1) A24=(ABLE*A18+BAKER*A13+CHARLY*A9)/(LAMBDA(ISOTOP)-LAMBDA(ISOTOP-4)) CHAINT 82
100 A25=(ABLE*A19+BAKER*A14+CHARLY*A10)/(LAMBDA(ISOTOP)-LAMBDA(ISOTOP-3)) CHAINT 83
A26=(ABLE*A20+BAKER*A15)/(LAMBDA(ISOTOP)-LAMBDA(ISOTOP-2)) CHAINT 84
A27=ABLE*A21/(LAMBDA(ISOTOP)-LAMBDA(ISOTOP-1)) CHAINT 85
105 A28=MASS(ISOTOP,POSIT)-A27-A26-A25-A24-A23-A22 CHAINT 86
MASS(ISOTOP,LOC)=A28*EXPO(7)+A27*EXPO(6)+A26*EXPO(5)+A25*EXPO(4)+ CHAINT 87
A24*EXPO(3)+A23*EXPO(2)+A22*EXPO(1) CHAINT 88
GO TO 9 CHAINT 89
110 8 J=0 CHAINT 90
9 CONTINUE CHAINT 91
GO TO 11 CHAINT 92
10 ISOTOP=ISOTOP+LIM CHAINT 93
11 CONTINUE CHAINT 94

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SUBROUTINE CHAINT 74/74 OPT=9 TRACE FTN 4.2+P386 12/10/74 1.32.55.

115	C	RETURN TO INDOSE	CHAINT	104
		RETURN	CHAINT	105
		END	CHAINT	106

	SUBROUTINE ORDOSE	ORDOSE	2
C	THIS ROUTINE CALCULATES THE DOSES TO SELECTED ORGANS DUE TO	ORDOSE	3
C	INHALATION OF SELECTED RADIONUCLIDES.	ORDOSE	4
5	C	ORDOSE	5
	COMMON NFP,NAC,NTOT,NOFNUC(96),SKIP(96),MASS(600,2),LAM9DA(600),	ORDOSE	6
	1 CHAINS,DKFRCT(45),3),REC(610),ML(600),ICL(600)	COMN	2
	COMMON MA(800),LAMBDA(1200),FMAT(1200),FAIR(1200),FZP(1200)	COMN	3
	COMMON IORG(1200),ENG(1200,3),F1(600),IDEG(600),IEND(600)	COMN	4
10	COMMON DOSTIM(10),D3,D4,D5,UPTIME(5),ISM(600)	COMN	5
	COMMON BRATE(5),R(10),X(10),SY(10),SZ(10),DIAM	COMN	6
	COMMON SUMDOS(50,10,10),TITLE2(8),NOT,KORG(20),TLH,TLN	COMN	7
	COMMON IPAGE,ORGAN(2,10)	COMN	8
	COMMON/NAM/ORGNAM(2,25)	COMN	9
15	REAL LAMBDA,LAMBDA,MASS	COMN	10
	INTEGER CHAINS,SKIP	COMN	11
	LOGICAL FISPRD,ACTPRD	COMN	12
	COMMON ACIPRU,FISPRU,KR,KUPT,DUY	COMN	13
	COMMON/COMO/ALAM,ALB(9),ALL(9),C(9),FT(9),NSOL,UMEJ1(9),	COMN	14
20	1 OMEJ2(9),OMEN2,T1,T2,VQNT1,VQLMT1,VQLUT1(8),OMASS(25)	COMO	2
	DIMENSION DOSE(10,10),ALLB(9,3),F(9,3),IBLNK(10,10),IFMT(17),	ORDOSE	3
	1 KFMT(21),ISPACE(10),TLH2(13)	ORDOSE	9
	DIMENSION ISOL(3)	ORDOSE	10
25	DATA ISOL/'(D) ','(M) ','(Y) '/	ORDOSE	11
	DATA (IFMT(I),I=1,6)/'(1H ','(1X,1X),'A3,I3','1PE9.2','2X)/	ORDOSE	12
	DATA (KFMT(I),I=1,21)/'(1H ','(4X,4X),'HRADI','2HO-',',2X, '	ORDOSE	13
	1'4HDOSE','1X,'4HTIME','/1H ','(4X,4X),'HNUCL','3HIOE','3X, '	ORDOSE	14
	2'4HIDAY','2HS),','(4X,4X),'46','45','/1H)'/	ORDOSE	15
	DATA ISPACE/'46','41','36','31','26','21','16','11','6','1'/	ORDOSE	16
30	DATA ((ALLB(I,J),I=1,9),J=1,3)/3*8.023E-4,4.012E-5,1.605E-5,2*0.,	ORDOSE	17
	12*1.605E-5,8.023E-4,2.036E-5,8.023E-4,4.012E-5,1.605E-7,8.023E-6,	ORDOSE	18
	23*1.605E-7,8.023E-4,2.036E-5,8.023E-4,4.012E-5,1.605E-8,8.023E-6,	ORDOSE	19
	32*1.605E-8,8.023E-9/	ORDOSE	20
	DATA ((F(I,J),I=1,9),J=1,3)/2*.5,.95,.05,.8,2*0..2,1..1,.9,2*.5,	ORDOSE	21
35	1.1,2*.4,.95,1..01,.99,.01,.99,.05,2*.4,.15,.9/	ORDOSE	22
	DATA (OMASS(I),I=1,25)/7.E4,4.3E4,307.,1700.,150.,7026.,1.E4,	ORDOSE	23
	11000.,20.,40.,8.,6100.,1500.,3.E4,20.,20.,70.,300.,2000.,250.,	ORDOSE	24
	21100.,135.,150.,0.,0./	ORDOSE	25
	C INITIALIZE VARIABLES	ORDOSE	26
40	DUMMY=0	ORDOSE	27
	ISO=0	ORDOSE	28
	ILINE=60	ORDOSE	29
	KRKUPT=KR + (KUPT-1)*10	ORDOSE	30
	SET I1 EQUAL TO THE CURRENT UPTAKE TIME	ORDOSE	31
45	I1=UPTIME(KUPT)	ORDOSE	32
	DO 90 I=1,NOT	ORDOSE	33
	TLH2(I)=(DOSTIM(I) + UPTIME(KUPT))/86400.	ORDOSE	34
90	CONTINUE	ORDOSE	35
	DO 100 I=1,10	ORDOSE	36
50	DO 100 J=1,10	ORDOSE	37
	DOSE(I,J)=0.	ORDOSE	38
	SUMDOS(KRKUPT,I,J)=0.0	ORDOSE	39
100	CONTINUE	ORDOSE	40
	C SET UP ORGAN NAMES FOR PRINT OUT	ORDOSE	41
55	LORG=0	ORDOSE	42
	DO 106 I=1,20	ORDOSE	43
	IF (KORG(I).EQ.0) GO TO 106	ORDOSE	44
		ORDOSE	45

		LORG=LORG+1	ORDOSE	46
		DO 105 J=1,2	ORDOSE	47
60	105	ORGAN(J,LORG)= ORGNAM(J,I)	ORDOSE	48
	106	CONTINUE	ORDOSE	49
		IFMT(LORG+7)=""	ORDOSE	50
		NORG=LORG	ORDOSE	51
		IF(IFISPRO) GO TO 200	ORDOSE	52
65		IF(ACTPRO) GO TO 700	ORDOSE	53
		WRITE(6,110) KUPT	ORDOSE	54
	110	FORMAT(1H0,"NO FISSION PRODUCTS OR ACTIVATION PRODUCTS SPECIFIED FOR	ORDOSE	55
		1R CASE",I3)	ORDOSE	56
		GO TO 3000	ORDOSE	57
70	C	LOOP ON NUCLIDE CHAINS	ORDOSE	58
	200	DO 600 ICH=1,CHAINS	ORDOSE	59
		LIM=NOFNUC(ICH)	ORDOSE	60
	C	BYPASS UNUSED CHAINS	ORDOSE	61
		IF(SKIP(ICH).EQ.3) GO TO 300	ORDOSE	62
75		ISO=ISO+LIM	ORDOSE	63
		GO TO 600	ORDOSE	64
	300	DO 500 K=1,LIM	ORDOSE	65
		ISO=ISO + 1	ORDOSE	66
		IF(MASS(ISO,2).LE.0.) GO TO 500	ORDOSE	67
80	C		ORDOSE	68
	C	IF DAUGHTER NUCLIDE SOLUBILITY INDEX HAS NOT BEEN SPECIFIED USE	ORDOSE	69
	C	THE SOLUBILITY INDEX OF THE PARENT	ORDOSE	70
		DO 302 J=1,K	ORDOSE	71
		IF(ICL(ISO).GT.0) GO TO 304	ORDOSE	72
85		ICL(ISO)=ICL(ISO-J)	ORDOSE	73
	302	CONTINUE	ORDOSE	74
		PRINT 303,ISO	ORDOSE	75
	303	FORMAT(1H0,"*** NO SOLUBILITY CLASS NUMBER FOR",I5," FOUND")	ORDOSE	76
	304	NSOL=ICL(ISO)	ORDOSE	77
90		L1=IBEG(ISO)	ORDOSE	78
		L2=IEND(ISO)	ORDOSE	79
		LORG=0	ORDOSE	80
		DO 305 L=1,9	ORDOSE	81
		ALL(L)=ALLB(L,NSOL) + LAMBOA(ISO)	ORDOSE	82
95		FT(L)=F(L,NSOL)	ORDOSE	83
		ARG=ALL(L)*T1	ORDOSE	84
		OHEJ1(L)=1. - EXP(-ARG)	ORDOSE	85
		IF(ARG.GT.1000.) OHEJ1(L)=1.	ORDOSE	86
		ALB(L)=ALLB(L,NSOL)	ORDOSE	87
100	305	CONTINUE	ORDOSE	88
		C(1)=ALB(1)*FT(1)*D3/ALL(1)	ORDOSE	89
		C(2)=ALB(2)*FT(2)*D3/ALL(2)*F1(ISO)	ORDOSE	90
		C(3)=ALB(3)*FT(3)*D4/ALL(3)	ORDOSE	91
		C(4)=ALB(4)*FT(4)*D4/ALL(4)*F1(ISO)	ORDOSE	92
105		C(5)=ALB(5)*FT(5)*D5/ALL(5)	ORDOSE	93
		C(6)=ALB(6)*FT(6)*D5/ALL(6)*F1(ISO)	ORDOSE	94
		C(7)=ALB(7)*FT(7)*D5/ALL(7)*F1(ISO)	ORDOSE	95
		C(8)=ALB(8)*ALB(9)*FT(8)*FT(9)*D5/ALL(8)	ORDOSE	96
		C(9)=C(8)/ALB(9)	ORDOSE	97
110		VQLMT1=QLMT1(DUMMY)	ORDOSE	98
		DO 306 J=1,8	ORDOSE	99
	306	VQLUT1(J)=QLUNT1(J)	ORDOSE	100
	C	ORGAN LOOP	ORDOSE	101
		DO 400 MORG=1,20	ORDOSE	102

115	C	TEST FOR ORGAN INCLUSION	ORDOSE= 113
		IF(KORG(MORG).EQ.0) GO TO 400	ORDOSE= 114
	C	TEST FOR ORGAN-NUCLIDE COMBINATION	ORDOSE= 115
		LORG=LORG + 1	ORDOSE= 116
		IF(L1.EQ.0.OR.L2.EQ.0) GO TO 312	ORDOSE= 117
120		DO 310 I0=L1,L2	ORDOSE= 118
		KRN=I0	ORDOSE= 119
		IF(IJRG(I0).EQ.MORG) GO TO 320	ORDOSE= 120
		310 CONTINUE	ORDOSE= 121
	C	NO ORGAN DATA AVAILABLE FOR THIS NUCLIDE - BLANK THE DOSE	ORDOSE= 122
125		312 DO 315 ITM=1,NDT	ORDOSE= 123
		IBLNK(ITM,LORG)=1	ORDOSE= 124
		315 DOSE(ITM,LORG)=' '	ORDOSE= 125
		GO TO 400	ORDOSE= 126
		320 ALAMO=LAMBIO(I0) +LAMBDA(ISO)	ORDOSE= 127
130		UM=N1=1. - EXP(-ALAMO*T1)	ORDOSE= 128
		VONT1=ONT1(KRN)	ORDOSE= 129
		D1=0.	ORDOSE= 130
	C	LOOP OVER DOSE TIMES	ORDOSE= 131
		DO 395 ITM=1,NDT	ORDOSE= 132
135		D2=0.	ORDOSE= 133
		T2=DOSTIM(ITM)	ORDOSE= 134
	C	ZERO OUT DOSE TIME FOR PRINT OUT IF LESS THAN 1 SEC	ORDOSE= 135
		IF(T2.LT.1.) T2=0.	ORDOSE= 136
		IBLNK(ITM,LORG)=0	ORDOSE= 137
140		IF(ITM.GT.1.AND.T2.LE.0.) GO TO 345	ORDOSE= 138
		OMEN2=1. - EXP(-ALAMO*T2)	ORDOSE= 139
		DO 375 J=1,9	ORDOSE= 140
		ARG=ALL(J)*T2	ORDOSE= 141
		OHEJ2(J)=1. - EXP(-ARG)	ORDOSE= 142
145		IF(ARG.GT.1000.) OHEJ2(J)=1.	ORDOSE= 143
		325 CONTINUE	ORDOSE= 144
		330 IF(ITM.GT.1) GO TO 340	ORDOSE= 145
		D1=ORDOT1(KRN,MORG)	ORDOSE= 146
		340 IF(T2.LE.0.) GO TO 345	ORDOSE= 147
150		D2=ORDOT2(KRN,MORG,ISO)	ORDOSE= 148
		345 IF(MORG.EQ.8) GO TO 360	ORDOSE= 149
		DOSE(ITM,LORG)=MASS(ISO,2) * LAMBDA(ISO) * (D1 + D2)	ORDOSE= 150
		GO TO 385	ORDOSE= 151
	C	ADD LUNG DOSE DUE TO PRIMARY DEPOSITION	ORDOSE= 152
155		360 SUM2=0.	ORDOSE= 153
		IF(ITM.GT.1) GO TO 370	ORDOSE= 154
		SUM1=0.	ORDOSE= 155
		DO 365 J=5,8	ORDOSE= 156
		365 SUM1=SUM1 + 5.92E-4*ENG(KRN,NSUL)/OMASS(MORG)*FT(J)*O5/ALL(J)	ORDOSE= 157
160		1*(T1 - OHEJ1(J)/ALL(J))	ORDOSE= 158
		370 IF(T2.LE.0.) GO TO 380	ORDOSE= 159
		DO 375 J=5,8	ORDOSE= 160
		375 SUM2=SUM2 + 5.92E-4*ENG(KRN,NSUL)/OMASS(MORG)*FT(J)*O5/ALL(J)**2	ORDOSE= 161
		1*OHEJ1(J)*OHEJ2(J)	ORDOSE= 162
165		380 DOSE(ITM,LORG)=MASS(ISO,2)*LAMBDA(ISO)*(SUM1+SUM2+D1+D2)	ORDOSE= 163
		385 SUMDOS(KRKUPT,ITM,LORG)=SUMDOS(KRKUPT,ITM,LORG) + DOSE(ITM,LORG)	ORDOSE= 164
	C	END OF TIME LOOP	ORDOSE= 165
		395 CONTINUE	ORDOSE= 166
	C	END OF ORGAN LOOP	ORDOSE= 167
170		400 CONTINUE	ORDOSE= 168
	C	SET UP TO WRITE DATA FOR ONE NUCLIDE	ORDOSE= 169

		IF (ILINE + NOT + 1.LE.60) GO TO 420	DRDOSE	160
	C	WRITE NEW PAGE HEADER	DRDOSE	161
		WRITE(6,410) IPAGE,DCY,TLH,TLN	DRDOSE	162
175	410	FORMAT(1H1,"PAGE",I5,31X,"DOSE TO SELECTED ORGANS OF INTEREST VIA 1 INHALATION",32X,A9/1H,49X,"TASK GROUP ON LUNG DYNAMICS MODEL" 21H,46X,"DURATION OF INHALATION",1PE9.2,1X,A6/1H,56X,"ORGAN DOSE 3 IN REM"//)	DRDOSE	163
		KFMT(2)=ISFACE(NORG)	DRDOSE	164
		KFMT(1)=KFMT(2)	DRDOSE	165
180		WRITE(6,KFMT) ((ORGAN(M,N),M=1,2),N=1,NORG)	DRDOSE	166
		IFMT(2)=KFMT(2)	DRDOSE	167
		IPAGE=IPAGE+1	DRDOSE	168
		ILINE=10	DRDOSE	169
185	420	DO 450 ITH=1,NDT	DRDOSE	170
		DO 430 I=1,NORG	DRDOSE	171
		IFMT(I+6)="E11.3"	DRDOSE	172
	430	IF (IBLNK(ITH,I).EQ.1) IFMT(I+6)="5X,A6"	DRDOSE	173
		IF (ITH.GT.1) GO TO 440	DRDOSE	174
190		IFMT(1)="(1H0,"	DRDOSE	175
		IFMT(4)="A3,I3,"	DRDOSE	176
		WRITE(6,IFMT) REC(ISO),ML(ISO),TLH2(1),(DOSE(1,I),I=1,NORG)	DRDOSE	177
		GO TO 450	DRDOSE	178
	440	IFMT(1)="(1H,"	DRDOSE	179
195		IFMT(4)="A6,"	DRDOSE	180
		MARK=ISOL(NSOL)	DRDOSE	181
		IF (ITH.GT.2) MARK=" "	DRDOSE	182
		WRITE(6,IFMT) MARK,TLH2(ITH),(DOSE(ITH,I),I=1,NORG)	DRDOSE	183
	450	CONTINUE	DRDOSE	184
200		ILINE=ILINE+NOT+1	DRDOSE	185
	C	END OF FP CHAIN LOOP	DRDOSE	186
	500	CONTINUE	DRDOSE	187
	C	END OF FP LOOP	DRDOSE	188
	650	CONTINUE	DRDOSE	189
205		IF (ACTPRO) GO TO 700	DRDOSE	190
		GO TO 2000	DRDOSE	191
	C	DO LOOP OVER ACTIVATION PRODUCTS	DRDOSE	192
	700	ISTART=NFP+1	DRDOSE	193
210		DO 1000 IS=ISTART,NTOT	DRDOSE	194
		IF (MASS(IS,2).LE.0.) GO TO 1000	DRDOSE	195
		L1=IBEG(IS)	DRDOSE	196
		L2=IEND(IS)	DRDOSE	197
		NSGL=ICL(IS)	DRDOSE	198
		LORG=0	DRDOSE	199
215		DO 705 L=1,9	DRDOSE	200
		ALL(L)=ALLB(L,NSOL) + LAMBDA(IS)	DRDOSE	201
		FT(L)=F(L,NSOL)	DRDOSE	202
		ARG=ALL(L)*T1	DRDOSE	203
		OMEJ1(L)=1. - EXP(-ARG)	DRDOSE	204
220		IF (ARG.GT.1000.) OMEJ1(L)=1.	DRDOSE	205
		ALB(L)=ALLB(L,NSOL)	DRDOSE	206
	705	CONTINUE	DRDOSE	207
		C(1)=ALB(1)*FT(1)*O3/ALL(1)	DRDOSE	208
		C(2)=ALB(2)*FT(2)*O3/ALL(2)*F1(IS)	DRDOSE	209
225		C(3)=ALB(3)*FT(3)*O4/ALL(3)	DRDOSE	210
		C(4)=ALB(4)*FT(4)*O4/ALL(4)*F1(IS)	DRDOSE	211
		C(5)=ALB(5)*FT(5)*O5/ALL(5)	DRDOSE	212
		C(6)=ALB(6)*FT(6)*O5/ALL(6)*F1(IS)	DRDOSE	213
			DRDOSE	214
			DRDOSE	215
			DRDOSE	216

		C(7)=ALB(7)*FT(7)*D5/ALL(7)*F1(15)	ORDOSE	217
230		C(8)=ALB(8)*ALB(9)*FT(8)*FT(9)*D5/ALL(8)	ORDOSE	218
		C(9)=C(8)/ALB(9)	ORDOSE	219
		VQLMT1=QLMT1(DDUMY)	ORDOSE	220
		DO 706 J=1,8	ORDOSE	221
	706	VQLUT1(J)=QLUNT1(J)	ORDOSE	222
235	C	ORGAN LOOP	ORDOSE	223
		DO 900 MORG=1,20	ORDOSE	224
	C	TEST FOR ORGAN INCLUSION	ORDOSE	225
		IF(KORG(MORG).EQ.0) GO TO 900	ORDOSE	226
	C	TEST FOR ORGAN-NUCLIDE COMBINATION DATA	ORDOSE	227
240		LORG=LORG + 1	ORDOSE	228
		DO 910 IO=L1,L2	ORDOSE	229
		KRN=IO	ORDOSE	230
		IF(IORG(IO).EQ.MORG) GO TO 820	ORDOSE	231
	810	CONTINUE	ORDOSE	232
245	C	NO ORGAN DATA FOR THIS NUCLIDE - BLANK DOSE ARRAY	ORDOSE	233
		DO 915 ITM=1,NOT	ORDOSE	234
		IBLNK(ITM,LORG)=1	ORDOSE	235
	815	DOSE(ITM,LORG)=-	ORDOSE	236
		GO TO 900	ORDOSE	237
250	820	ALAMO= LAMBIO(IO) + LAMBOA(15)	ORDOSE	238
		OMEN1=1. - EXP(-ALAMO*T1)	ORDOSE	239
		VQNT1=QNT1(KRN)	ORDOSE	240
		D1=0.	ORDOSE	241
	C	THE FOLLOWING TWO CARDS WERE ADDED TO ALLOW CHANGING THE MASS OF	9/4/74	3
255	C	THE LUNG TO 570 GRAMS IF A ALPHA EMITTER DOSE IS BEING COMPUTED.	9/4/74	4
		OMCF=1.	9/4/74	5
		IF(MORG.EQ.8.AND.ENG(KRN,NSOL).GT.10.) OMCF=1.754	9/4/74	6
	C	LOOP OVER DOSE TIMES	ORDOSE	242
		DO 895 ITM=1,NOT	ORDOSE	243
260		D2=J.	ORDOSE	244
		T2=JOSTIM(ITM)	ORDOSE	245
	C	ZERO OUT DOSE TIME FOR PRINT OUT IF LESS THAN 1 SEC	ORDOSE	246
		IF(T2.LT.1.) T2=J.	ORDOSE	247
		IBLNK(ITM,LORG)=0	ORDOSE	248
265		IF(ITM.GT.1.AND.T2.LE.0.) GO TO 845	ORDOSE	249
		OMEN2=1. - EXP(-ALAMO*T2)	ORDOSE	250
		DO 825 J=1,9	ORDOSE	251
		ARG=ALL(J)*T2	ORDOSE	252
		OMEJ2(J)=1. - EXP(-ARG)	ORDOSE	253
270		IF(ARG.GT.1000.) OMEJ2(J)=1.	ORDOSE	254
	825	CONTINUE	ORDOSE	255
	830	IF(ITM.GT.1) GO TO 840	ORDOSE	256
		D1=ORDOT1(KRN,MORG)	ORDOSE	257
	840	IF(T2.LE.0) GO TO 845	ORDOSE	258
275		D2=GRDOT2(KRN,MORG,15)	ORDOSE	259
	845	IF(MORG.EQ.8) GO TO 860	ORDOSE	260
		DOSE(ITM,LORG)= MASS(15,2)*LAMBOA(15) * (D1 + D2)	ORDOSE	261
		GO TO 885	ORDOSE	262
280	C	ADD LUNG DOSE FOR THAT DEPOSITED DIRECTLY	ORDOSE	263
	860	SUM2=0.	ORDOSE	264
		IF(ITM.GT.1) GO TO 870	ORDOSE	265
		SUM1=0.	ORDOSE	266
		DO 365 J=5,8	ORDOSE	267
	365	SUM1=SUM1 + 5.92E-4*ENG(KRN,NSOL)/CMASS(MORG)*FT(J)*D5/ALL(J)	ORDOSE	268
285		1*(T1 - OMEJ1(J)/ALL(J)) * OMCF	9/4/74	7

	870	IF(T2.LE.0.) GO TO 880	ORDOSE	270
		DO 875 J=5,8	ORDOSE	271
	875	SUM2=SUM2 + 5.92E-4*ENG(KRN,NSOL)/OMASS(NORG)*FT(J)*D5/ALL(J)**2	ORDOSE	272
		1*OMEJ1(J)*CMEJ2(J) + UMGF	9/4/74	8
290	880	DOSE(ITM,LORG)=MASS(IS,2)*LAMBDA(IS)*(SUM1+SUM2+D1+D2)	ORDOSE	274
	885	SUMDOS(KRKUPT,ITM,LORG)=SUMDOS(KRKUPT,ITM,LORG) + DOSE(ITM,LORG)	ORDOSE	275
	C	END OF TIME LOOP	ORDOSE	276
	895	CONTINUE	ORDOSE	277
	C	END OF ORGAN LOOP	ORDOSE	278
295	900	CONTINUE	ORDOSE	279
	C	SET UP TO WRITE DATA FOR ONE NUCLIDE	ORDOSE	280
		IF (ILINE + NDT + 1.LE.60) GO TO 920	ORDOSE	281
	C	WRITE NEW PAGE HEADER	ORDOSE	282
		WRITE(6,410) IPAGE,DOY,TLH,TLN	ORDOSE	283
300		KFMT(2)=ISFACE(NORG)	ORDOSE	284
		KFMT(11)=KFMT(2)	ORDOSE	285
		WRITE(6,KFMT) ((ORGAN(M,N),M=1,2),N=1,NORG)	ORDOSE	286
		IFMT(2)=KFMT(2)	ORDOSE	287
		IFMT(NORG+7)=""	ORDOSE	288
305		IPAGE=IPAGE+1	ORDOSE	289
		ILINE=10	ORDOSE	290
	920	DO 950 ITM=1,NDT	ORDOSE	291
		DO 930 I=1,NORG	ORDOSE	292
		IFMT(I+6)="",E11.3"	ORDOSE	293
310	930	IF (IBLNK(ITM,I).EQ.1) IFMT(I+6)="5X,A6"	ORDOSE	294
		IF (ITM.GT.1) GO TO 940	ORDOSE	295
		IFMT(1)="1H0,"	ORDOSE	296
		IFMT(4)="A3,I3,"	ORDOSE	297
315		WRITE(6,IFMT) REC(IS),ML(IS),TLH2(1),(DOSE(1,I),I=1,NORG)	ORDOSE	298
		GO TO 950	ORDOSE	299
	940	IFMT(1)="1H,"	ORDOSE	300
		IFMT(4)="A6,"	ORDOSE	301
		MARK=ISOL(NSOL)	ORDOSE	302
		IF (ITM.GT.2.OR.NSOL.EQ.0) MARK=" "	ORDOSE	303
320		WRITE(6,IFMT) MARK,TLH2(ITM),(DOSE(ITM,I),I=1,NORG)	ORDOSE	304
	950	CONTINUE	ORDOSE	305
		ILINE=ILINE+NDT+1	ORDOSE	306
	C	END OF ACTIVATION PRODUCT LOOP	ORDOSE	307
	1000	CONTINUE	ORDOSE	308
325	C	SUMMARY PAGE OUTPUT	ORDOSE	309
	2000	WRITE(6,2001) IPAGE,KUPT,KR,TITLE2,DOY	ORDOSE	310
	2001	FORMAT(1H1,"PAGE",I5,1X,"(",I2,"",I2,"")",8X,8A10,17X,A9)	5/30/74	1
		WRITE(6,2010) TLH,TLN	ORDOSE	312
330	2010	FORMAT(1H0,40X,"DOSE TO SELECTED ORGANS OF INTEREST VIA INHALATION	ORDOSE	313
		1"/1H,48X,"TASK GROUP ON LUNG DYNAMICS MODEL"/1H,46X,	ORDOSE	314
		2"DURATION OF INHALATION",1F9.2,1X,A6/1H,57X,"ORGAN DOSE IN REM"/	ORDOSE	315
		1/)	ORDOSE	316
		WRITE(6,KFMT) ((ORGAN(I,J),I=1,2),J=1,NORG)	ORDOSE	317
		IFMT(1)="1H0,"	ORDOSE	318
335		IFMT(4)="A6,"	ORDOSE	319
		HALL="ALL "	ORDOSE	320
		DO 2015 I=1,NORG	ORDOSE	321
	2015	IFMT(I+6)="",E11.2"	ORDOSE	322
		IPAGE=IPAGE+1	ORDOSE	323
340		DO 2020 I=1,NDT	ORDOSE	324
		WRITE(6,IFMT) HALL,TLH2(I),(SUMDOS(KRKUPT,I,J),J=1,NORG)	ORDOSE	325
	2020	CONTINUE	ORDOSE	326

SUBROUTINE ORDOSE 7474 OPT=0 TRACE

FIN 4.2+390

12/16/74 14.38.52.

3000 RETURN
END

ORDOSE 327
ORDOSE 328

		FUNCTION ORDOT1(KRN,MORG)	ORDOT1	2
	C	ORDOT1 CALCULATES THE NORMALIZED DOSE TO AN ORGAN DURING THE TIME	ORDOT1	3
	C	OF UPTAKE T1.	ORDOT1	4
5	C	"C" ARE THE COEF. FOR THE RESPIRATORY TRACT CLEARANCE	ORDOT1	5
	C	"ALAMO" IS THE EFFECTIVE REMOVAL CONSTANT FOR THE ORGAN	ORDOT1	6
	C	"ALL" ARE THE EFFECTIVE REMOVAL CONSTANTS FOR THE LUNG COMPARTMENT	ORDOT1	7
	C	"OMEN1" IS (1 - EXP(-ALAMO * T1))	ORDOT1	8
	C	"T1" IS THE UPTAKE TIME	ORDOT1	9
10	C	"NSOL" IS THE SOLUBILITY INDEX	ORDOT1	10
	C	"F2P" IS THE BLOOD TO ORGAN FRACTION	ORDOT1	11
	C	"FT" ARE THE RESPIRATORY TRACT CLEARANCE PATHWAY FRACTIONS	ORDOT1	12
	C	"ALB" ARE THE BIOLOGICAL REMOVAL CONSTANTS FOR THE LUNG	ORDOT1	13
	C	"ENG" IS THE EFFECTIVE ENERGY FOR THE NUCLIDE IN THE ORGAN	ORDOT1	14
15	C	"OMASS" IS THE ORGAN MASS	ORDOT1	15
	C	"D3,D4,D5" ARE THE FRACTION INHALED DEPOSITION FACTORS	ORDOT1	16
	C	"OMEJ1" IS (1 - EXP(-ALL*T1))	ORDOT1	17
		COMMON NFP,NAC,NTOT,NCFNUC(96),SKIP(96),MASS(600,2),LAMBDA(630),	COMN	18
		1CHAINS,DKFRCT(43),3),REC(600),ML(600),ICL(600)	COMN	2
20		COMMON HA(600),LAMBDA(1200),FWAT(1200),FAIR(1200),F2P(1200)	COMN	3
		COMMON IORG(1200),ENG(1200,3),F1(600),IBEG(600),IEND(600)	COMN	4
		COMMON DOSTIM(10),D3,D4,D5,UPTIME(5),ISM(600)	COMN	5
		COMMON BRATE(5),R(10),X(10),SY(10),SZ(10),OIAM	COMN	6
		COMMON SUNDUS(50,10,10),TITLE2(8),NDT,KORG(20),TLH,TLN	COMN	7
25		COMMON IPAGE,ORGAN(2,10)	COMN	8
		COMMON/NAM/ORGNAH(2,25)	COMN	9
		REAL LAMBDA,LAMBDA,MASS	COMN	10
		INTEGER CHAINS,SKIP	COMN	11
		LOGICAL FISPRD,ACTPRD	COMN	12
30		COMMON ACTPRG,FISPRD,KR,KUPT,DOY	COMN	13
		COMMON/COMO/ALAMO,ALB(9),ALL(9),C(9),FT(9),NSOL,OMEJ1(9),	COMN	14
		1OMEJ2(9),OMEN1,OMEN2,T1,T2,VQNT1,VQLMT1,VQLUT1(8),OMASS(25)	COMO	2
		FIRST=0.	COMO	3
		W=T1/ALAMO	ORDOT1	21
35		Y=OMEN1/ALAMO	ORDOT1	22
		Z=Y/ALAMO	ORDOT1	23
		DO 30 I=1,7	ORDOT1	24
		IF(NSOL.EQ.1.AND.(I.EQ.6.OR.I.EQ.7)) GO TO 30	ORDOT1	25
		FIRST=FIRST +C(I)*(W-Z-(Y -OMEJ1(I))/ALL(I))/(ALL(I)-ALAMO)	ORDOT1	26
40	30	CONTINUE	ORDOT1	27
		SECND=C(6)*(W-Z-Y/(ALL(9)-ALAMO)+OMEJ1(9)/ALL(9))/(ALL(9)-	ORDOT1	28
		ALAMO)/ALL(9)	ORDOT1	29
		IF(NSOL.LE.2) GO TO 40	ORDOT1	30
	C	ALL(8).NE.ALL(9)	ORDOT1	31
45		THIRD=C(8)/(ALL(8)-ALL(9))*(Y-OMEJ1(9)/ALL(9))/(ALL(9)-ALAMO)	ORDOT1	32
		1-Y-OMEJ1(8)/ALL(8))/(ALL(8)-ALAMO)	ORDOT1	33
		GO TO 100	ORDOT1	34
	C	ALL(8).EQ.ALL(9)	ORDOT1	35
		THIRD=C(9)/(ALL(9)-ALAMO)**2 * (OMEJ1(9)/ALL(9)-Y+(ALL(9)-ALAMO)	ORDOT1	36
50	40	1/ALL(9)**2 * (1. -EXP(-ALL(9)*T1)*(ALL(9)*T1 +1.))	ORDOT1	37
	100	ORDOT1=5.92E-4*ENG(KRN,NSOL)*F2P(KRN)/OMASS(MORG)*(FIRST+SECND	ORDOT1	38
		1 + THIRD)	ORDOT1	39
		END	ORDOT1	40
			ORDOT1	41

	C	FUNCTION ORDOT2(KRN,MOKG,ISO)	ORDOT2	2
	C	ODOT2 CALCULATES THE NORMALIZED DOSE TO AN ORGAN DURING THE TIME	ORDOT2	3
	C	FOLLOWING TERMINATION OF UPTAKE T2	ORDOT2	4
5	C	"C" ARE THE COEF. FOR RESPIRATORY TRACT CLEARANCE	ORDOT2	5
	C	"ALAMO" IS THE EFFECTIVE REMOVAL CONSTANT FOR THE ORGAN	ORDOT2	6
	C	"ALL" ARE THE EFFECTIVE REMOVAL CONSTANTS FOR THE LUNG COMPARTMENTS	ORDOT2	7
	C	"OMEN1" AND "OMEN2" ARE (1 - EXP(-ALAMO*T1)) AND (1 - EXP(-ALAMO*T2))	ORDOT2	8
	C	"OMEJ1" AND "OMEJ2" ARE (1 - EXP(-ALL*I1)) AND (1 - EXP(-ALL*I2))	ORDOT2	9
10	C	"T1" IS THE UPTAKE TIME	ORDOT2	10
	C	"T2" IS THE DOSE TIME FOLLOWING UPTAKE	ORDOT2	11
	C	"NSOL" IS THE SOLUBILITY INDEX	ORDOT2	12
	C	"F2P" IS THE BLOOD TO ORGAN INDEX	ORDOT2	13
	C	"FT" ARE THE RESPIRATORY TRACT CLEARANCE PATHWAY FRACTIONS	ORDOT2	14
15	C	"ALD" ARE THE BIOLOGICAL REMOVAL CONSTANTS FOR THE LUNG	ORDOT2	15
	C		ORDOT2	16
		COMMON HFP,NAC,NTOT,NOFNUG(96),SKIP(96),MASS(600,2),LAMBDA(630),	COMN	2
		1CHAINS,OKFRCT(450,3),REC(600),HL(600),ICL(600)	COMN	3
20		COMMON MA(600),LAMBID(1200),FWAT(1200),FAIR(1200),F2P(1200)	COMN	4
		COMMON IORG(1200),ENG(1200,3),F1(600),IBEG(600),IEND(600)	COMN	5
		COMMON DOSTIM(10),J3,0,05,UPTIME(5),ISM(600)	COMN	6
		COMMON BRATE(5),R(10),X(10),SY(10),SZ(10),DIAM	COMN	7
		COMMON SUMDOS(50,10,10),TITLE2(8),NOT,KORG(20),TLH,TLN	COMN	8
25		COMMON IPAGE,ORGAN(2,10)	COMN	9
		COMMON/NAM/ORGAN(2,25)	COMN	10
		REAL LAMBIC,LAMBDA,MASS	COMN	11
		INTEGER CHAINS,SKIP	COMN	12
		LOGICAL FISPRD,ACTPRD	COMN	13
		COMMON ACTFRD,FISPRD,KR,KUPT,DOY	COMN	14
30		COMMON/COMG/ALAMO,ALB(9),ALL(9),C(9),FT(9),NSOL,OMEJ1(9),	COMO	2
		OMEJ2(9),OMEN1,OMEN2,T1,T2,VQNT1,VQLMT1,VQLUT1(8),OMASS(25)	COMO	3
		FIRST=OMEN2/ALAMO * VQNT1	ORDOT2	20
		SECOND=0.	ORDOT2	21
		DO 30 I=1,7	ORDOT2	22
35		FGIB=1.	ORDOT2	23
		IF(MOD(I,2).EQ.0.OR.I.EQ.7) FGIB=F1(ISO)	ORDOT2	24
		IF(NSOL.EQ.1.AND.(I.EQ.6.OR.I.EQ.7)) GO TO 30	ORDOT2	25
		SECOND=SECOND + ALB(I) / (ALL(I)-ALAMO)*(OMEN2/ALAMO-OMEJ2(I)/	ORDOT2	26
40		1ALL(I))*VQLUT1(I)*FGIB	ORDOT2	27
	30	CONTINUE	ORDOT2	28
		IF(NSOL.LE.2) GO TO 40	ORDOT2	29
		THIRD=ALB(8) * ALB(9) * FT(9)*VQLUT1(8)/(ALL(8)-ALL(9)) *	ORDOT2	30
		1((OMEN2 /ALAMO-OMEJ2(9))/ALL(9))/(ALL(9)-ALAMO)	ORDOT2	31
45		2-(OMEN2 /ALAMO - OMEJ2(9)/ALL(8))/(ALL(8)-ALAMO)	ORDOT2	32
		GO TO 100	ORDOT2	33
	40	THIRD=ALB(8) * ALB(9) * FT(9)*VQLUT1(8)/(ALL(9)-ALAMO)*((OMEN2/ALAMO	ORDOT2	34
		1-OMEJ2(9)/ALL(9))/(ALL(9)-ALAMO) - (1 - EXP(-ALL(9)*T2)*(ALL(9)*T2	ORDOT2	35
		2 + 1.))/ALL(9)**2)	ORDOT2	36
50	100	FOURTH=ALB(9) * VQLMT1/(ALAMO-ALL(9))*(OMEJ2(9)/ALL(9)	ORDOT2	37
		1 - OMEJ2/ALAMO)	ORDOT2	38
		ORDOT2=5.92E-4 * ENG(KRN,NSOL)/OMASS(MOKG)*(FIRST + F2P(KRN)	ORDOT2	39
		1 * (SECOND + THIRD + FOURTH))	ORDOT2	40
		RETURN	ORDOT2	41
		END	ORDOT2	42

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FUNCTION QNT1(KRN)
C / THIS FUNCTION CALCULATES THE QUANTITY OF A NUCLIDE PRESENT IN AN OR
C FOLLOWING CONTINUOUS UPTAKE AT A UNIT RATE FOR A TIME T1 USING
C THE NEW ICRP LUNG MODEL.
5 C
C "T1" IS THE UPTAKE TIME
C "C" ARE THE COEF. FOR THE RESPIRATORY TRACT CLEARANCE
C "ALAMO" IS THE EFFECTIVE REMOVAL CONSTANT FOR THE ORGAN
C "ALL" ARE THE EFFECTIVE REMOVAL CONSTANTS FOR THE LUNG COMPARTMENTS
10 C "OMEN" IS (1 - EXP(-ALAMO*T1))
C "OMEJ" ARE (1 - EXP(-ALL(J)*T1))
C "NSOL" IS THE SOLUBILITY INDEX
C "F2P" IS THE BLOOD TO ORGAN FRACTION
15 C
COMMON NFP,NAC,NTOT,NOFNUC(96),SKIP(96),MASS(600,2),LAMBDA(600),
1 CHAINS,OKFRCT(450,3),REC(600),HL(600),ICL(600)
COMMON MA(600),LAMBDA(1200),FNAT(1200),FAIR(1200),F2P(1200)
COMMON IORG(1200),ENG(1200,3),F1(600),IDEG(600),IEND(600)
COMMON JOSTIM(10),D3,D4,D5,UPTIME(5),ISM(600)
20 COMMON BRATE(5),R(10),X(10),SY(10),SZ(10),DIAM
COMMON SUMDDS(50,10,10),TITLE2(8),NDT,KORG(20),TLH,TLN
COMMON IPAGE,ORGAN(2,10)
COMMON/NAM/ORGNAM(2,25)
REAL LAMBIC,LAMBDA,MASS
25 INTEGER CHAINS,SKIP
LOGICAL FISPRD,ACTPRD
COMMON ACTPRD,FISPRD,KR,KUPT,DJY
COMMON/COMO/ALAMO,ALB(9),ALL(9),C(9),FT(9),NSOL,OMEJ1(9),
30 OMEJ2(9),OMEN1,OMEN2,T1,T2,VQNT1,VQLHT1,VQLUT1(8),OMASS(25)
SUM=0.
DO 50 I=1,7
IF(NSOL.EQ.1.AND.(I.EQ.6.OR.I.EQ.7)) GO TO 50
SUM=SUM + C(I) * (OMEN1/ALAMO - (OMEJ1(I) - OMEN1)/(ALL(I)-ALAMO))
50 CONTINUE
35 C
SUM=SUM + C(8)/ALL(9) * (OMEN1/ALAMO - (OMEJ1(9) - OMEN1)/(ALL(9)-ALAMO))
IF (NSOL.LE.2) GO TO 70
C THIS CASE IS FOR WHEN ALL(8) NOT EQUAL TO ALL(9)
SUM=SUM - C(8)/(ALL(8)-ALL(9)) * ((OMEJ1(9) - OMEN1)/(ALL(9)-ALAMO)
40 1 - (OMEJ1(8) - OMEN1)/(ALL(8)-ALAMO))
GO TO 100
C THIS CASE IS FOR ALL(8)=ALL(9)
70 SUM=SUM - C(8)/(ALL(9)-ALAMO)**2 * (EXP(-ALAMO*T1)
1 - EXP(-ALL(9)*T1) * ((ALL(9)-ALAMO)*T1 + 1.))
45 100 QNT1=SUM * F2P(KRN)
RETURN
END

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QNT1      2
QNT1      3
QNT1      4
QNT1      5
QNT1      6
QNT1      7
QNT1      8
QNT1      9
QNT1     10
QNT1     11
QNT1     12
QNT1     13
QNT1     14
QNT1     15
COMN      2
COMN      3
COMN      4
COMN      5
COMN      6
COMN      7
COMN      8
COMN      9
COMN     10
COMN     11
COMN     12
COMN     13
COMN     14
COMO      2
COMO      3
QNT1     18
QNT1     19
QNT1     20
QNT1     21
QNT1     22
QNT1     23
QNT1     24
QNT1     25
QNT1     26
QNT1     27
QNT1     28
QNT1     29
QNT1     30
QNT1     31
QNT1     32
QNT1     33
QNT1     34
QNT1     35

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	C	FUNCTION QLUNT1(I)	QLUNT1	2
	C		QLUNT1	3
	C	THIS FUNCTION CALCULATES THE QUANTITY OF A NUCLIDE PRESENT IN	QLUNT1	4
	C	EACH OF THE LUNG SUB COMPARTMENTS FOLLOWING CONTINUOUS INHALATION	QLUNT1	5
5	C	AT A UNIT RATE FOR A TIME T1 USING THE NEW ICRP LUNG MODEL.	QLUNT1	6
	C		QLUNT1	7
	C	"FT" ARE THE RESPIRATORY TFACT CLEARANCE PATHWAY FRACTIONS	QLUNT1	8
	C	"D3,D4,D5" ARE THE LUNG DEPOSITION FACTORS	QLUNT1	9
	C	"OMEJ1" IS (1 - EXP(-ALL*T1))	QLUNT1	10
10	C	"ALL" ARE THE EFFECTIVE REMOVAL CONSTANTS FOR THE LUNG COMPARTMENTS	QLUNT1	11
	C		QLUNT1	12
		COMMON NFP,NAC,NTDT,NOFNUC(96),SKIP(96),MASS(600,2),LAMBDA(600),	COMMON	2
		1CHAINS,OKFRCT(450,3),REC(600),ML(600),ICL(600)	COMMON	3
		COMMON MA(600),LAMBDA(1200),FWAT(1200),FAIR(1200),F2P(1200)	COMMON	4
15		COMMON IORG(1200),ENG(1200,3),F1(600),I3EG(600),IEND(600)	COMMON	5
		COMMON DOSTI(10),D3,D4,D5,UPTIME(5),IS4(600)	COMMON	6
		COMMON BRATE(5),R(10),X(10),SY(10),SZ(10),DIAM	COMMON	7
		COMMON SUMDJS(50,10,10),TITLE2(8),NOT,KORG(20),TLH,ILN	COMMON	8
		COMMON IFACE,ORGAN(2,10)	COMMON	9
20		COMMON/NAH/ORGNAH(2,25)	COMMON	10
		REAL LAMBDA,LAMBDA,MASS	COMMON	11
		INTEGER CHAINS,SKIP	COMMON	12
		LOGICAL FISPRD,ACTPRD	COMMON	13
		COMMON ACTPRD,FISPRD,KR,KUPT,DOY	COMMON	14
25		COMMON/COMO/ALAMO,ALB(9),ALL(9),C(9),FT(9),NSOL,OMEJ1(9),	COMMON	2
		1OMEJ2(9),OMEN1,OMEN2,T1,T2,VQNT1,VQLUT1(8),OMASS(25)	COMMON	3
		GO TO (10,10,20,20,30,30,30,30), I	QLUNT1	15
10		QLUNT1= FT(I) * D3 * OMEJ1(I)/ALL(I)	QLUNT1	16
		GO TO 100	QLUNT1	17
30		20 QLUNT1= FT(I) * D4 * OMEJ1(I)/ALL(I)	QLUNT1	18
		GO TO 100	QLUNT1	19
		30 QLUNT1= FT(I) * D5 * OMEJ1(I)/ALL(I)	QLUNT1	20
100		RETURN	QLUNT1	21
		END	QLUNT1	22

	FUNCTION QLMIT1(DUMMY)	QLMIT1	
		2	
		3	
		4	
5	THIS FUNCTION CALCULATES THE QUANTITY OF NUCLIDE PRESENT IN THE LYMPH, THAT IS REMOVABLE, FOLLOWING CONTINUOUS UPTAKE AT A UNIT RATE FOR A TIME T1 USING THE NEW LUNG MODEL.	5	
		6	
		7	
	"ALL" ARE THE REMOVAL CONSTANTS FOR THE RESPIRATORY TRACT	8	
	"OMEJ1" ARE (1 - EXP(-ALL(J)*T1))	9	
	"C" ARE THE COEF. FOR THE RESPIRATORY TRACT CLEARANCE	10	
10	"T1" IS THE UPTAKE TIME	11	
	"NSOL" IS THE SOLUBILITY INDEX	12	
	COMMON NFP, NAC, NTOT, NOFNUC(96), SKIP(96), MASS(600,2), LAMBDA(600),	COMN	2
	1CHAINS, DKFRCT(450,3), REC(600), ML(600), ICL(600)	COMN	3
	COMMON MA(600), LAMBIO(1200), FMAT(1200), FAIR(1200), F2P(1200)	COMN	4
15	COMMON IORG(1200), ENG(1200,3), F1(600), IBEG(600), IEND(600)	COMN	5
	COMMON DOSTIM(10), D3, D4, D5, UPTIME(5), ISM(600)	COMN	6
	COMMON BRATE(5), R(10), X(10), SY(10), SZ(10), DIAM	COMN	7
	COMMON SUMDOS(50,10,10), TITLE2(8), NOT, KORG(20), TLH, TLN	COMN	8
	COMMON IPAGE, ORGAN(2,10)	COMN	9
20	COMMON/NAH/ORGNAH(2,25)	COMN	10
	REAL LAMBIO, LAMBDA, MASS	COMN	11
	INTEGER CHAINS, SKIP	COMN	12
	LOGICAL FISPRD, ACTPRD	COMN	13
	COMMON ACTPRD, FISPRD, KR, KUPT, DOY	COMN	14
25	COMMON/COMO/ALAMO, ALB(9), ALL(9), C(9), FT(9), NSOL, OMEJ1(9),	COMO	2
	1OMEJ2(9), OMEN1, OMEN2, T1, T2, VQNT1, VQLMT1, VQLUT1(8), OMASS(25)	COMO	3
	DUMY=DUMMY	QLMIT1	15
	FIRST= OMEJ1(9)/ALL(9)	QLMIT1	16
	IF(NSOL.LE.2) GO TO 50	QLMIT1	17
30	C ALL8.NE.ALL9	QLMIT1	18
	QLMIT1= C(9)*(FIRST - (OMEJ1(8) - OMEJ1(9))/(ALL(8)-ALL(9)))	QLMIT1	19
	GO TO 100	QLMIT1	20
	C ALL8.EQ.ALL9	QLMIT1	21
	50 QLMIT1= C(9)* (FIRST - T1 * EXP(-ALL(9)*T1))	QLMIT1	22
35	100 RETURN	QLMIT1	23
	END	QLMIT1	24

	SUBROUTINE UNIT(DTLH,DTLN,NUPT,UPTIME)	UNIT	2
C	UNIT CONVERTS TIME IN SECONDS TO MORE CONVENIENT UNITS	UNIT	3
	DIMENSION DTLH(5),DTLN(5),UPTIME(5)	UNIT	4
	DC 100 I=1,NUPT	UNIT	5
5	IF (UPTIME(I).GT.60.) GO TO 10	UNIT	6
	DTLH(I)=UPTIME(I)	UNIT	7
	DTLN(I)="SEC"	UNIT	8
	GO TO 100	UNIT	9
	10 IF (UPTIME(I).GT.3600.) GO TO 20	UNIT	10
10	DTLH(I)=UPTIME(I)/60.	UNIT	11
	DTLN(I)="MIN"	UNIT	12
	GO TO 100	UNIT	13
	20 IF (UPTIME(I).GT.86400.) GO TO 30	UNIT	14
	DTLH(I)=UPTIME(I)/3600.	UNIT	15
15	DTLN(I)="HOURS"	UNIT	16
	GO TO 100	UNIT	17
	30 IF (UPTIME(I).GT.3.1536E7) GO TO 40	UNIT	18
	DTLH(I)=UPTIME(I)/86400.	UNIT	19
	DTLN(I)="DAYS"	UNIT	20
20	GO TO 100	UNIT	21
	40 DTLH(I)=UPTIME(I)/3.1536E7	UNIT	22
	DTLN(I)="YEARS"	UNIT	23
	100 CONTINUE	UNIT	24
	RETURN	UNIT	25
25	END	UNIT	26

APPENDIX F

INPUT PREPARATION

APPENDIX F

INPUT PREPARATION

The input data required for the calculations falls into three general categories: (1) Program control variables, (2) Parameters associated with atmospheric diffusion and dispersion and (3) Parameters associated with the organ dose calculation. The majority of the data required by the code is contained in two data libraries that are accessed by the code (see Appendix D) thus eliminating much time consuming preparation of input.

Program control variables consist of parameters to control the reading of the data libraries, the optional printing of some input and intermediate data and the type of atmospheric dispersion calculation to be performed, if any. The atmospheric dispersion calculation can be by-passed by setting MET = 0; in this case, the values entered for the radionuclide activity become the total quantity inhaled rather than the quantity released to the atmosphere.

The remainder of this appendix describes in detail, the input variables, their use and the manner in which they are input to the program.

DATA CARD DESCRIPTIONS

CARD 1 - Title card

<u>COLUMN</u>	<u>FORMAT</u>	<u>VARIABLE</u>	<u>USE</u>
1-72	12A6	TITLE 2	words entered here are reprinted as a title on the summary page. Each case requires a title card or blank card.

CARDS 2 to N - Namelist Data Cards (one or more namelist cards).

The minimum data for each case consists of one title card (card 1) one or more cards using the Namelist format (cards 2 to N). The first Namelist card (card 2) must be blank in Column 1, \$INPUT in columns 2-7,

followed by at least one blank, followed by data items. The data items are separated by a comma, and the last data item must be followed by a \$. The data items must have one of the three following forms:

1. Variable name = constant, where the variable name may be either subscripted or not.
2. Array name = set of constants (separated by commas). The number of constants must not exceed the number of elements in the array and they must be in the same order as the array is in storage, i.e., the first subscript changes most rapidly.
3. Subscripted variable = set of constants (separated by commas). This form results in the set of constants being placed in consecutive array elements, starting with the element designated by the subscripted variable. Again, the number of elements in the array between the given element in the array and the last element in the array.

The namelist variables retain their values throughout the execution of the program and need not be respecified unless a change is wished. The namelist input variables are shown in the following list:

TABLE OF NAMELIST VARIABLES

I. PROGRAM CONTROL VARIABLES

<u>VARIABLE</u>	<u>ARRAY SIZE</u>	<u>USE</u>	<u>VALUE UNITS</u>	<u>VALUES</u>
NEXT	(1)	Next Case Options	None	1 to 4
IPR	(1)	Controls printing of u_z values (no or yes)	None	0 or 1
IPU	(1)	Controls punching of E/Q values on cards (no or yes)	None	0 or 1
IQP	(1)	Controls printing of E/Q values (no or yes)	None	0 or 1

<u>VARIABLE</u>	<u>ARRAY SIZE</u>	<u>DESCRIPTION</u>	<u>VALUE UNITS</u>	<u>VALUES</u>
MET	(1)	Selects atmospheric dispersion model to be used	None	0 to 10

II. ATMOSPHERIC DISPERSION VARIABLES

A. GENERAL

<u>VARIABLE</u>	<u>ARRAY SIZE</u>	<u>DESCRIPTION</u>	<u>VALUE UNITS</u>	<u>VALUES</u>
H	(1)	Release height	meters	any
R(K)	(10)	Radial distance to receptor	meters	any
UBAR	(1)	Wind speed	m/sec	any

B. SIMPSON-FUQUAY (MET=1)

<u>VARIABLE</u>	<u>ARRAY SIZE</u>	<u>DESCRIPTION</u>	<u>VALUE UNITS</u>	<u>VALUES</u>
STUB(J)	(5)	Meteorological parameters, ($\sigma_{\theta} \bar{u}$)	radian-m/sec	any
AH	(1)	Meteorological parameter, a	m ²	any
BH	(1)	Meteorological parameter, b	m ² /sec	any
CH	(1)	Meteorological parameter, c	m ² /sec	any
DH	(1)	Meteorological parameter, d	sec	any
KSQD	(1)	Meteorological parameter, k ²	sec ⁻²	any

C. SUTTON (MET=2)

<u>VARIABLE</u>	<u>ARRAY SIZE</u>	<u>DESCRIPTION</u>	<u>VALUE UNITS</u>	<u>VALUES</u>
CY	(1)	Sutton's lateral diffusion coefficient, C _y	None	any

<u>VARIABLE</u>	<u>ARRAY SIZE</u>	<u>DESCRIPTION</u>	<u>VALUE UNITS</u>	<u>VALUES</u>
CZ	(1)	Sutton's vertical diffusion coefficient, C_z	None	any
EN	(1)	Sutton's power coefficient, n	None	any

D. ANY OTHER MODEL DESCRIBED BY σ_z AND σ_y (MET=3)

<u>VARIABLE</u>	<u>ARRAY SIZE</u>	<u>DESCRIPTION</u>	<u>VALUE UNITS</u>	<u>VALUES</u>
SIGY (K,I)	(10,5)	Standard deviation of lateral plume growth, σ_y at the Kth distance for the Ith inhalation uptake time.	meters	any
SIGZ (K,I)	10,5)	Standard deviation of vertical plume growth, σ_z at the Kth distance for the Ith inhalation uptake time.	meters	any

E. PASQUILL (MET=4 through 9)

No special atmospheric dispersion variables required.

III. ORGAN DOSE VARIABLES

<u>VARIABLE</u>	<u>ARRAY SIZE</u>	<u>DESCRIPTION</u>	<u>VALUE UNITS</u>	<u>VALUES</u>
D3	(1)	Fraction of aerosol inhaled deposited in nasal passages	None	any
D4	(1)	Fraction of aerosol inhaled deposited in trachea and bronchia	None	any
D5	(1)	Fraction of aerosol inhaled deposited in pulmonary region	None	any

<u>VARIABLE</u>	<u>ARRAY SIZE</u>	<u>DESCRIPTION</u>	<u>VALUE UNITS</u>	<u>VALUES</u>
DIAM	(1)	Mass median aerodynamic diameter of inhaled material	μm	any
BRATE(I)	(5)	Ventilation rate of receptor during the Ith inhalation uptake time	cm^3/sec	any
UPTIME(I)	(5)	The length of time during which inhalation uptake takes place	sec	any
DOSTIM(J)	(10)	The length of time following the termination of inhalation uptake for which the dose is to be calculated	sec	any
ORGANS(L)	(10)	Selects the organ's for which the dose is to be calculated (see the organ list for numerical code)	None	1 to 18 (integer)

NEXT

Next is a variable controlling the dose calculations that instructs the program what to do after the namelist data has been read. It controls library reading and quantitative isotopic input data. There are 4 options corresponding to NEXT - 1,2,3,4.

NEXT = 1 Reads in libraries, zeros out isotopic quantities and reads in a set of isotope quantities.

NEXT = 2 Zeros out existing isotopes and reads in a new set of isotope quantities.

- NEXT = 3 Isotopic quantities are kept, other variables may be changed.
- NEXT = 4 End of job. No further calculations are done. Provides for normal exit.

Any of the namelist variables may be changed when NEXT = 1, 2 or 3. NEXT must be specified as 1 in the first case so the libraries will be read.

MET

MET is a variable used to select the atmospheric dispersion model to calculate the E/Q values. There are 11 options available corresponding to MET = 0,1,2,3,...,10.

- MET = 0 By-passes atmospheric dispersion portion of the code. Only program control and organ dose namelist variables are required.
- MET = 1 Simpson-Fuquay Model for stable atmospheric conditions. The unique namelist variables required for this case are shown under II.B.
- MET = 2 Sutton's Model for neutral and unstable atmospheric conditions. The unique namelist variables required for this case are shown under II.C.
- MET = 3 Any other situation where the values of σ_y and σ_z are known. The unique namelist variables required for this case are shown under II.D.
- MET = 4 Pasquill's Model for atmospheric conditions "A" through to 9 "F". MET = 4 corresponds to condition "A", etc. No unique namelist variables are required for this case. The values for σ_y and σ_z are interpolated from built-in data.
- MET = 10 Input of E/Q values directly. No unique namelist variables are required for this case.

CARD N+1 - Number of Radionuclides Card

<u>COLUMN</u>	<u>FORMAT</u>	<u>VARIABLE</u>	<u>USE</u>
1-3	I3	NUMBER	Controls the number of radionuclides (maximum of 600) to be read from subsequent input cards.

CARDS N+2 to M - Radionuclide Data Cards

<u>COLUMN</u>	<u>FORMAT</u>	<u>VARIABLE</u>	<u>USE</u>
1-3	A3	NAME	The name of an element selected from the Isotope Data Library
4-6	I3	M	The mass number of the selected radionuclide
7-8	I2	ISL	The solubility class of the nuclide (1=D, 2=W, 3=Y)
9-20	E12.0	QUANTITY	The total quantity of this radionuclide released to the atmosphere or if MET=0 the total inhaled.

The format shown may be repeated 4 times on each card starting in columns 21, 41, and 61 for additional nuclides.

CARDS M+1 to K - E/Q Value Input Cards (used only with MET=10)

<u>COLUMN</u>	<u>FORMAT</u>	<u>VARIABLE</u>	<u>USE</u>
1-70	10E7.3	EQQ	E/Q values for up to ten downwind distances

One of these cards is required for each uptake time specified only if MET=10 is used.

ORGAN LIST

<u>NUMERICAL CODE</u>	<u>ORGAN</u>
1	TOTAL BODY
2	BODY WATER
3	KIDNEYS
4	LIVER
5	SPLEEN
6	BONE
7	FAT
8	LUNGS
9	ADRENALS
10	TESTES
11	OVARIES
12	SKIN
13	BRAIN
14	MUSCLE
15	PROSTATE
16	THYROID
17	PANCREAS
18	HEART

APPENDIX G

SAMPLE PROBLEMS

APPENDIX G

SAMPLE PROBLEMS

To illustrate the use of the computer code, two sample problems are presented here. One using the direct inhalation option and one using the Simpson-Fuquay stable atmospheric diffusion model option, to simulate the inhalation of radionuclides released to the atmosphere.

The first problem includes calculation of the effective dose to ten different organs or tissues from the chronic inhalation of a mixture of fission products for five dose times following the termination of uptake.

The second problem will make use of the atmospheric diffusion model built into the program to calculate the dose to four organs at two dose times following passage of the cloud produced by the short term release to the atmosphere of a mixture of transuranic nuclides.

Input for the first problem is:

<u>Name List Parameter</u>	<u>Value</u>	<u>Remarks</u>
NEXT	1	Program control variables.
MET	0	See Appendix F.
D3	0.30	Respiratory tract disposition factors for one micrometer particles.
D4	0.08	
D5	0.25	
UPTIME	3.156×10^7	Time of uptakes in seconds (one year)
DOSTIM	1	Dose times in seconds (zero time not allowed by data checking portion of program).
	3.156×10^7	
	6.312×10^8	
	1.578×10^8	
	3.156×10^8	

Name List Parameter	Value	Remarks
ORGANS		Organs Considered:
	1	Whole Body
	3	Kidneys
	4	Liver
	5	Spleen
	6	Bone
	8	Lung
	10	Testes
	14	Muscle
	16	Thyroid
	18	Heart

Other Parameters	Value	Remarks
NUMBER	9	Number of radionuclides to be read from succeeding cards.
M, NAME ISL, QUANTY	⁹⁰ Sr 2 1x10 ⁻⁶	Radionuclide identifiers, solubility index and quantity in curies inhaled.
	⁹⁰ Y 2 1x10 ⁻⁶	
	⁹⁵ Zr 3 2x10 ⁻⁶	
	⁹⁵ Nb 3 1x10 ⁻⁶	
	¹⁰⁶ Ru 3 5x10 ⁻⁶	
	¹³¹ I 1 1x10 ⁻⁵	
	¹³⁷ Cs 1 1x10 ⁻⁶	
	¹⁴⁴ Ce 3 2x10 ⁻⁶	
	¹⁴⁴ Pr 3 2x10 ⁻⁶	

If both problems are submitted in the same deck, input for the second problem is as follows:

Name List Parameter	Value	Remarks
NEXT	2	Program control variables.
MET	1	General atmospheric dispersion variables.
H	70	Height in meters.

Name List Parameter	Value	Remarks
R	100,200,300,500,1000,2000, 3000,5000,10000,20000	Downwind distances in meters.
UBAR	1.	Windspeed in meters/sec. Simpson-Fuquay Model Variables.
STUB	0.04	$(\sigma_{\theta} \bar{u})$ radian-meters/sec
AH	97	a, m ²
BH	0.33	b, m ² /sec
CH	13	c, m ² /sec
DH	230	d, sec
KSQD	0.00025	k ² , sec ⁻²
D3 D4 D5	Same as preceding case--no need to respecify	Respiratory tract deposition factors.
BRATE	350	Ventilation rate of receptor in cm ³ /sec
UPTIME	3600	Uptake time in seconds (one hour)
DOSTIM	3.156x10 ⁷	Dosetimes in seconds (one year and 50 years)
ORGANS		Organs Considered:
	3	Kidneys
	4	Liver
	6	Bone
	8	Lung

<u>Other Parameters</u>	<u>Value</u>		<u>Remarks</u>
NUMBER		3	Number of radionuclides to be read from succeeding cards.
M, NAME	²³⁸ Pu	3	Radionuclide identifiers, solubility index and quantity in curies released to the atmosphere.
ISL, QUANTY	²³⁹ Pu	3	
	²⁴⁰ Pu	3	
		5×10^{-4}	
		4×10^{-2}	
		1×10^{-2}	

Shown in Table G.1 is a listing of the input data card stream prepared from the preceding information.

Output reports generated by the code for the two sample problems are shown following the input listing.

SAMPLE INPUT

SAMPLE PROBLEM NUMBER ONE CHRONIC INHALATION
\$INPUT NEXT=1,MET=0,D3=.3,D4=.08,D5=.25,UPTIME=3.156E+7,4*0.,
DOSTIM=1.,3.156E+7,6.312E+7,1.578E+8,3.156E+8,5*0.,ORGANS=1,3,4,5,
6,8,10,14,16,18\$END

SR 90 2 1.F-6Y 90 2 1.E-6ZR 95 3 2.E-6NB 95 3 1.E-6
RU 106 3 5.E-6I 131 1 1.E-5CS 137 1 1.E-6CE 144 3 2.E-6
PR 144 3 2.F-6

SAMPLE PROBLEM NUMBER TWO ACUTE INHALATION
\$INPUT NEXT=2,MET=1,H=70.0,R=100.,200.,300.,500.,1000.,2000.,3000.,
5000.,10000.,20000.,UBAR=1.,STUR=.04,4*0.,AH=97.,BH=.330,CH=13.,DH=230.,
KSQD=2.5F-4,PRATF=350.,4*0.,UPTIME=3600.,4*0.,IPR=1,IQP=1,
ORGANS=3,4,6,8,6*0,DOSTIM=3.156E+7,1.578E+9,8*0.\$

3
PU 238 3 5.E-4PU 239 3 4.E-2PU 240 3 1.E-2
END OF RUN
\$INPUT NEXT=4\$

SAMPLE OUTPUT

PAGE 1

***** INPUT DATA *****

11/08/74

ORGAN DATA LIBRARY VERSION 10 SEPTEMBER 24, 1974

NUCLIDE RELEASE IN CURIES

SR 90 2 1,0000E-06 Y 90 2 1,0000E-06 ZR 95 3 2,0000E-06 NB 95 3 1,0000E-06 RU 106 3 5,0000E-06 I 131 1 1,0000E-05
CS 137 1 1,0000E-06 CE 144 3 2,0000E-06 PR 144 3 2,0000E-06

UBAR = 0.0

BRATE = 0. 0. 0. 0. 0.

NET = 0

H0 = 0 METERS

1 UPTAKE TIMES, SECONDS 3.156E+07

5 DOSE TIMES, SECONDS 1,000E+00 3,156E+07 6,312E+07 1,578E+08 3,156E+08

DOSE TO SELECTED ORGANS OF INTEREST VIA INHALATION
TASK GROUP ON LUNG DYNAMICS MODEL
DURATION OF INHALATION 1.00E+00 YEARS
ORGAN DOSE IN REM

11/08/78

RADIO-NUCLIDE	DOSE TIME (DAYS)	TOTAL BODY	KIDNEYS	LIVER	SPLEEN	BONE	LUNGS	TESTES	MUSCLE	THYROID	HEART
BR 90	3.65E+02	3.769E+02				1.416E+01	1.318E+01				
	(M) 7.31E+02	1.141E+01				4.295E+01	1.612E+01				
	1.10E+03	1.876E+01				7.081E+01	1.614E+01				
	2.19E+03	3.898E+01				1.483E+00	1.614E+01				
	4.02E+03	6.734E+01				2.993E+00	1.614E+01				
Y 90	3.65E+02	2.048E+04				7.592E+03	1.233E+02				
	(M) 7.31E+02	2.070E+04				7.677E+03	1.241E+02				
	1.10E+03	2.070E+04				7.677E+03	1.241E+02				
	2.19E+03	2.070E+04				7.677E+03	1.241E+02				
	4.02E+03	2.070E+04				7.677E+03	1.241E+02				
Zn 95	3.65E+02	5.132E+04	1.049E+03	7.393E+04	6.292E+04	1.944E+03	5.612E+01				
	(Y) 7.31E+02	7.211E+04	1.499E+03	1.026E+03	8.997E+04	2.784E+03	7.193E+01				
	1.10E+03	7.274E+04	1.515E+03	1.033E+03	9.090E+04	2.814E+03	7.212E+01				
	2.19E+03	7.275E+04	1.515E+03	1.034E+03	9.093E+04	2.815E+03	7.212E+01				
	4.02E+03	7.275E+04	1.515E+03	1.034E+03	9.093E+04	2.815E+03	7.212E+01				
NR 95	3.65E+02	7.174E+05	1.313E+04	1.360E+04	8.325E+05	1.996E+04	8.384E+02				
	(Y) 7.31E+02	8.523E+05	1.560E+04	1.617E+04	9.540E+05	2.375E+04	9.600E+02				
	1.10E+03	8.525E+05	1.560E+04	1.617E+04	9.541E+05	2.375E+04	9.600E+02				
	2.19E+03	8.525E+05	1.560E+04	1.617E+04	9.541E+05	2.375E+04	9.600E+02				
	4.02E+03	8.525E+05	1.560E+04	1.617E+04	9.541E+05	2.375E+04	9.600E+02				
RU 106	3.65E+02	1.085E+03	1.665E+02			8.312E+03	6.879E+00				
	(Y) 7.31E+02	1.329E+03	2.002E+02			1.053E+02	1.353E+01				
	1.10E+03	1.428E+03	2.168E+02			1.134E+02	1.553E+01				
	2.19E+03	1.496E+03	2.249E+02			1.190E+02	1.637E+01				
	4.02E+03	1.501E+03	2.254E+02			1.194E+02	1.640E+01				
I 131	3.65E+02	2.093E+02					2.600E+02			1.149E+01	
	(D) 7.31E+02	2.159E+02					2.605E+02			1.185E+01	
	1.10E+03	2.159E+02					2.605E+02			1.185E+01	
	2.19E+03	2.159E+02					2.605E+02			1.185E+01	
	4.02E+03	2.159E+02					2.605E+02			1.185E+01	
Ca 137	3.65E+02	1.994E+02	2.001E+02	4.879E+02	3.585E+02	2.780E+02	8.047E+03			2.733E+02	
	(D) 7.31E+02	2.706E+02	2.398E+02	6.869E+02	5.423E+02	4.763E+02	1.110E+02			4.683E+02	
	1.10E+03	2.725E+02	2.399E+02	6.998E+02	5.558E+02	5.080E+02	1.159E+02			4.996E+02	
	2.19E+03	2.725E+02	2.399E+02	7.006E+02	5.564E+02	5.141E+02	1.168E+02			5.055E+02	
	4.02E+03	2.725E+02	2.399E+02	7.006E+02	5.564E+02	5.141E+02	1.168E+02			5.055E+02	
Ce 144	3.65E+02	1.363E+03	6.266E+03	1.251E+02			2.096E+02			2.426E+00	
	(Y) 7.31E+02	3.542E+03	1.653E+02	3.055E+02			5.883E+02			4.545E+00	
	1.10E+03	4.788E+03	2.234E+02	3.940E+02			8.310E+02			5.070E+00	
	2.19E+03	5.706E+03	2.663E+02	4.501E+02			1.042E+01			5.240E+00	
	4.02E+03	5.741E+03	2.688E+02	4.528E+02			1.046E+01			5.243E+00	

DOSE TO SELECTED ORGANS OF INTEREST VIA INHALATION
 TASK GROUP ON LUNG DYNAMICS MODEL
 DURATION OF INHALATION 1,00E+00 YEARS
 ORGAN DOSE IN REM

RADIO-NUCLIDE	DOSE TIME (DAYS)	TOTAL BODY	KIDNEYS	LIVER	SPLEEN	BONE	LUNGS	TESTES	MUSCLE	THYROID	HEART
ALL	3.05E+02	8.18E+02	4.41E+02	6.01E+02	3.66E+02	2.08E+01	1.01E+01	0.	2.73E+02	1.15E+01	0.
ALL	7.31E+02	1.09E+01	6.22E+02	1.00E+01	5.52E+02	5.57E+01	1.91E+01	0.	4.68E+02	1.19E+01	0.
ALL	1.10E+03	2.44E+01	6.95E+02	1.11E+01	5.66E+02	8.60E+01	2.16E+01	0.	5.00E+02	1.19E+01	0.
ALL	2.19E+03	4.47E+01	7.48E+02	1.16E+01	5.67E+02	1.66E+00	2.26E+01	0.	5.06E+02	1.19E+01	0.
ALL	4.02E+03	7.31E+01	7.51E+02	1.17E+01	5.67E+02	2.77E+00	2.27E+01	0.	5.06E+02	1.19E+01	0.

ORGAN DATA LIBRARY VERSION 1B SEPTEMBER 24, 1974

NUCLIDE RELEASE IN CURIES

PU-238-3-5.0000E-04 PU-239-3-4.0000E-02 PU-240-3-1.0000E-02

UBAR = 1.0
 BRATE = 350. 0. 0. 0. 0.
 MET = 1
 HE 70 METERS
 AMS 97.0
 BWR 33000
 CWR 13.0
 D= 230.0
 KSD = 0.000250
 STUB(1) = 0.040
 DIAM = 0.000 MICROMETERS
 D3 = 300
 D4 = 0.080
 D5 = 250

10 DISTANCES, METERS 100. 200. 300. 500. 1000. 2000. 3000. 5000. 10000. 20000.
 1 UPTAKE TIMES, SECONDS 5.600E+03
 2 DOSE TIMES, SECONDS 5.150E+07 1.578E+09

SIGMA Z VALUES

AVERAGE WIND SPEED, M/SEC	DISTANCE FROM RELEASE POINT METERS									
	100.	200.	300.	500.	1000.	2000.	3000.	5000.	10000.	20000.
1.0	11.	13.	14.	16.	21.	28.	33.	42.	50.	82.

SIGMA Y VALUES

AVERAGE WIND SPEED, M/SEC	DISTANCE FROM RELEASE POINT METERS									
	100.	200.	300.	500.	1000.	2000.	3000.	5000.	10000.	20000.
1.0	4.	8.	12.	20.	39.	76.	112.	179.	323.	546.

E/O VALUES

UPTAK TIMES / DISTANCES
(SECONDS) (METERS)

	100.	200.	300.	500.	1000.	2000.	3000.	5000.	10000.	20000.	
	3.400E+03	1.380E+11	9.286E+10	7.112E+09	8.004E+08	1.271E+06	5.957E+06	9.055E+06	1.069E+05	8.216E+06	8.917E+06

DOSE TO SELECTED ORGANS OF INTEREST VIA INHALATION
 TASK GROUP ON LUNG DYNAMICS MODEL
 DURATION OF INHALATION 6.00E+01 MIN
 ORGAN DOSE IN REM

11/05/74

RADIO-NUCLIDE	DOSE-TIME (DAYS)	KIDNEYS	LIVER	BONE	LUNGS
PU 238	3.65E+02	1.209E-12	4.781E-12	5.729E-12	5.307E-10
(Y)	1.83E+04	2.704E-10	6.942E-10	1.307E-09	1.317E-09
PU 239	3.65E+02	9.033E-11	3.572E-10	4.439E-10	3.962E-08
(Y)	1.83E+04	2.635E-08	7.954E-08	1.222E-07	9.948E-08
PU 240	3.65E+02	2.258E-11	6.930E-11	1.110E-10	9.904E-09
(Y)	1.83E+04	6.975E-09	1.935E-08	3.048E-08	2.487E-08

DOSE TO SELECTED ORGANS OF INTEREST VIA INHALATION
TASK GROUP ON LUNG DYNAMICS MODEL
DURATION OF INHALATION $6.00E+01$ MIN
ORGAN DOSE IN REM

RADIO- NUCLIDE	DOSE TIME (DAYS)	KIDNEYS	LIVER	BONE	LUNGS
ALL	$3.65E+02$	$1.14E+10$	$4.51E+10$	$5.61E+10$	$5.01E+08$
ALL	$1.83E+04$	$3.07E+08$	$1.00E+07$	$1.54E+07$	$1.26E+07$

DOSE TO SELECTED ORGANS OF INTEREST VIA INHALATION
TASK GROUP ON LUNG DYNAMICS MODEL
DURATION OF INHALATION 6,00E+01 MIN
ORGAN DOSE IN REM

11/05/74

RADIO-NUCLIDE	DOSE TIME (DAYS)	KIDNEYS	LIVER	BONE	LUNGS
PU 238	3,65E+02	8,137E-11	3,218E-10	3,856E-10	3,572E-08
(Y)	1,83E+04	1,820E-08	6,018E-08	8,795E-08	8,867E-08
PU 239	3,65E+02	6,079E-09	2,404E-08	2,988E-08	2,666E-06
(Y)	1,83E+04	1,639E-06	5,353E-06	8,222E-06	6,695E-06
PU 240	3,65E+02	1,520E-09	6,011E-09	7,469E-09	6,666E-07
(Y)	1,83E+04	4,089E-07	1,336E-06	2,051E-06	1,674E-06

DOSE TO SELECTED ORGANS OF INTEREST VIA INHALATION
TASK GROUP ON LUNG DYNAMICS MODEL
DURATION OF INHALATION $6.00E+01$ MIN
ORGAN DOSE IN REM

RADIO-NUCLIDE	DOSE TIME (DAYS)	KIDNEYS	LIVER	BONE	LUNGS
ALL	$3.65E+02$	$7.68E-09$	$3.04E-08$	$3.77E-08$	$3.37E-06$
ALL	$1.83E+04$	$2.07E-06$	$6.75E-06$	$1.04E-05$	$8.46E-06$

DOSE TO SELECTED ORGANS OF INTEREST VIA INHALATION
 TASK GROUP ON LUNG DYNAMICS MODEL
 DURATION OF INHALATION 6.00E+01 MIN
 ORGAN DOSE IN REM

11/05/74

RADIO-NUCLIDE	DOSE TIME (DAYS)	KIDNEYS	LIVER	BONE	LUNGS
PU 238	3.65E+02	6.231E-10	2.464E-09	2.953E-09	2.736E-07
(Y)	1.83E+04	1.304E-07	4.609E-07	6.735E-07	6.791E-07
PU 239	3.65E+02	4.656E-08	1.841E-07	2.288E-07	2.042E-05
(Y)	1.83E+04	1.255E-05	4.100E-05	6.297E-05	5.127E-05
PU 240	3.65E+02	1.164E-08	4.603E-08	5.720E-08	5.105E-06
(Y)	1.83E+04	3.132E-06	1.023E-05	1.571E-05	1.282E-05

DOSE TO SELECTED ORGANS OF INTEREST VIA INHALATION
TASK GROUP ON LUNG DYNAMICS MODEL
DURATION OF INHALATION 6.00E+01 MIN
ORGAN DOSE IN REM

RADIO-NUCLIDE	DOSE TIME (DAYS)	KIDNEYS	LIVER	BONE	LUNGS
ALL	3.65E+02	5.88E+08	2.35E+07	2.89E+07	2.58E+05
ALL	1.83E+04	1.58E+05	5.17E+05	7.94E+05	6.48E+05

DOSE TO SELECTED ORGANS OF INTEREST VIA INHALATION
 TASK GROUP ON LUNG DYNAMICS MODEL
 DURATION OF INHALATION 6.0E+01 MIN
 ORGAN DOSE IN REM

11/05/78

RADIO-NUCLIDE	DOSE-TIME (DAYS)	KIDNEYS	LIVER	BONE	LUNGS
PU 238	3.65E+02	7.574E-09	2.995E-08	3.589E-08	3.325E-06
(Y)	1.83E+04	1.694E-06	5.602E-06	8.186E-06	8.254E-06
PU 239	3.65E+02	5.659E-07	2.238E-06	2.781E-06	2.482E-04
(Y)	1.83E+04	1.523E-04	4.983E-04	7.654E-04	6.232E-04
PU 240	3.65E+02	1.415E-07	5.595E-07	6.952E-07	6.205E-05
(Y)	1.83E+04	3.806E-05	1.243E-04	1.910E-04	1.558E-04

DOSE TO SELECTED ORGANS OF INTEREST VIA INHALATION
TASK GROUP ON LUNG DYNAMICS MODEL
DURATION OF INHALATION $6.00E+03$ MIN
ORGAN DOSE IN REM

RADIO- NUCLIDE	DOSE TIME (DAYS)	KIDNEYS	LIVER	BONE	LUNGS
ALL	$3.65E+02$	$7.15E+07$	$2.83E+06$	$3.51E+06$	$3.14E+04$
ALL	$1.83E+04$	$1.92E+04$	$6.28E+04$	$9.64E+04$	$7.87E+04$

DOSE TO SELECTED ORGANS OF INTEREST VIA INHALATION
 TASK GROUP ON LUNG DYNAMICS MODEL
 DURATION OF INHALATION $6.00E+01$ MIN
 ORGAN DOSE IN REM

11/05/74

RADIO-NUCLIDE	DOSE TIME (DAYS)	KIDNEYS	LIVER	BONE	LUNGS
PU 238	$3.65E+02$	$1.113E-07$	$4.403E-07$	$5.276E-07$	$4.888E-05$
(Y)	$1.83E+04$	$2.491E-05$	$8.235E-05$	$1.203E-04$	$1.213E-04$
PU 239	$3.65E+02$	$8.319E-06$	$3.290E-05$	$4.088E-05$	$3.649E-03$
(Y)	$1.83E+04$	$2.243E-03$	$7.325E-03$	$1.125E-02$	$9.161E-03$
PU 240	$3.65E+02$	$2.080E-06$	$8.224E-06$	$1.022E-05$	$9.121E-04$
(Y)	$1.83E+04$	$5.595E-04$	$1.828E-03$	$2.807E-03$	$2.200E-03$

DOSE TO SELECTED ORGANS OF INTEREST VIA INHALATION
TASK GROUP ON LUNG DYNAMICS MODEL
DURATION OF INHALATION 6.00E+01 MIN
ORGAN DOSE IN REM

RADIO- NUCLIDE	DOSE TIME (DAYS)	KIDNEYS	LIVER	BONE	LUNGS
ALL	3.65E+02	1.05E-05	4.16E-05	5.16E-05	4.61E-03
ALL	1.83E+04	2.83E-03	9.24E-03	1.42E-02	1.16E-02

DOSE TO SELECTED ORGANS OF INTEREST VIA INHALATION
TASK GROUP ON LUNG DYNAMICS MODEL
DURATION OF INHALATION 6,00E+01 MIN
ORGAN DOSE IN REM

RADIO-NUCLIDE	DOSE-TIME (DAYS)	KIDNEYS	LIVER	BONE	LUNGS
PU 238	3.65E+02	5.219E+07	2.064E+06	2.473E+06	2.291E+04
(Y)	1.83E+04	1.168E+04	3.860E+04	5.641E+04	5.688E+04
PU 239	3.65E+02	3.900E+05	1.542E+04	1.916E+04	1.710E+02
(Y)	1.83E+04	1.051E+02	3.434E+02	5.274E+02	4.295E+02
PU 240	3.65E+02	9.749E+06	3.856E+05	4.791E+05	4.276E+03
(Y)	1.83E+04	2.623E+03	8.568E+03	1.316E+02	1.074E+02

DOSE TO SELECTED ORGANS OF INTEREST VIA INHALATION
TASK GROUP ON LUNG DYNAMICS MODEL
DURATION OF INHALATION 6.00E+01 MIN
ORGAN DOSE IN REM

RADIO-NUCLIDE	DOSE TIME (DAYS)	KIDNEYS	LIVER	BONE	LUNGS
ALL	3.65E+02	4.93E-05	1.95E-04	2.42E-04	2.16E-02
ALL	1.83E+04	1.33E-02	4.33E-02	6.65E-02	5.43E-02

DOSE TO SELECTED ORGANS OF INTEREST VIA INHALATION
TASK GROUP ON LUNG DYNAMICS MODEL
DURATION OF INHALATION 6,00E+01 MIN
ORGAN DOSE IN REM

11/05/74

RADIO-NUCLIDE	DOSE TIME (DAYS)	KIDNEYS	LIVER	BONE	LUNGS
PU 238	3.65E+02	7.934E+07	3.138E+06	3.760E+06	3.483E+04
(Y)	1.83E+04	1.775E+04	5.868E+04	8.576E+04	8.646E+04
PU 239	3.65E+02	5.928E+05	2.344E+04	2.913E+04	2.600E+02
(Y)	1.83E+04	1.598E+02	5.220E+02	8.018E+02	6.529E+02
PU 240	3.65E+02	1.482E+05	5.861E+05	7.283E+05	6.500E+03
(Y)	1.83E+04	3.987E+03	1.303E+02	2.000E+02	1.632E+02

DOSE TO SELECTED ORGANS OF INTEREST VIA INHALATION
TASK GROUP ON LUNG DYNAMICS MODEL
DURATION OF INHALATION 6.00E+01 MIN
ORGAN DOSE IN REM

RADIO-NUCLIDE	DOSE TIME (DAYS)	KIDNEYS	LIVER	BONE	LUNGS
ALL	3.65E+02	7.49E+05	2.96E+04	3.68E+04	3.28E+02
ALL	1.83E+04	2.01E+02	6.58E+02	1.01E+01	8.25E+02

DOSE TO SELECTED ORGANS OF INTEREST VIA INHALATION
 TASK GROUP ON LUNG DYNAMICS MODEL
 DURATION OF INHALATION 6.00E+01 MIN
 ORGAN DOSE IN REM

11/05/74

RADIO-NUCLIDE	DOSE-TIME (DAYS)	KIDNEYS	LIVER	BONE	LUNGS
PU 238	3.65E+02	9.191E-07	3.635E-06	4.355E-06	4.035E-04
(Y)	1.83E+04	2.056E-04	6.798E-04	9.934E-04	1.002E-03
PU 239	3.65E+02	6.867E-05	2.716E-04	3.375E-04	3.012E-02
(Y)	1.83E+04	1.851E-02	6.047E-02	9.288E-02	7.563E-02
PU 240	3.65E+02	1.717E-05	6.789E-05	8.437E-05	7.530E-03
(Y)	1.83E+04	4.610E-03	1.509E-02	2.317E-02	1.890E-02

DOSE TO SELECTED ORGANS OF INTEREST VIA INHALATION
TASK GROUP ON LUNG DYNAMICS MODEL
DURATION OF INHALATION $6.00E+01$ MIN
ORGAN DOSE IN REM

RADIO-NUCLIDE	DOSE TIME (DAYS)	KIDNEYS	LIVER	BONE	LUNGS
ALL	$3.65E+02$	$8.68E+05$	$3.43E+04$	$4.26E+04$	$3.81E+02$
ALL	$1.83E+04$	$2.33E+02$	$7.62E+02$	$1.17E+01$	$9.55E+02$

DOSE TO SELECTED ORGANS OF INTEREST VIA INHALATION
 TASK GROUP ON LUNG DYNAMICS MODEL
 DURATION OF INHALATION 6.00E+01 MIN
 ORGAN DOSE IN RPM

11/09/74

RADIO- NUCLIDE	DOSE TIME (DAYS)	KIDNEYS	LIVER	BONE	LUNGS
PU 238	3.65E+02	7.199E-07	2.847E-06	3.412E-06	3.161E-04
(Y)	1.83E+04	1.610E-04	5.328E-04	7.781E-04	7.845E-04
PU 239	3.65E+02	5.379E-05	2.127E-04	2.643E-04	2.359E-02
(Y)	1.83E+04	1.450E-02	4.737E-02	7.275E-02	5.924E-02
PU 240	3.65E+02	1.345E-05	5.318E-05	6.608E-05	5.898E-03
(Y)	1.83E+04	3.618E-03	1.182E-02	1.815E-02	1.481E-02

DOSE TO SELECTED ORGANS OF INTEREST VIA INHALATION
TASK GROUP ON LUNG DYNAMICS MODEL
DURATION OF INHALATION 6.00E+01 MIN
ORGAN DOSE IN REM

RADIO-NUCLIDE	DOSE TIME (DAYS)	KIDNEYS	LIVER	BONE	LUNGS
ALL	3.65E+02	6.80E-05	2.69E-04	3.34E-04	2.98E-02
ALL	1.83E+04	1.83E-02	5.97E-02	9.17E-02	7.48E-02

DOSE TO SELECTED ORGANS OF INTEREST VIA INHALATION
TASK GROUP ON LUNG DYNAMICS MODEL
DURATION OF INHALATION $6.00E+01$ MIN
ORGAN DOSE IN REM

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RADIO- NUCLIDE	DOSE-TIME (DAYS)	KIDNEYS	LIVER	BONE	LUNGS
PU 238	$3.65E+02$	$4.326E-07$	$1.711E-06$	$2.050E-06$	$1.899E-04$
(Y)	$1.83E+04$	$9.677E-05$	$3.200E-04$	$4.676E-04$	$4.714E-04$
PU 239	$3.65E+02$	$3.232E-05$	$1.278E-04$	$1.588E-04$	$1.418E-02$
(Y)	$1.83E+04$	$8.713E-03$	$2.846E-02$	$4.372E-02$	$3.560E-02$
PU 240	$3.65E+02$	$8.080E-06$	$3.196E-05$	$3.971E-05$	$3.544E-03$
(Y)	$1.83E+04$	$2.174E-03$	$7.102E-03$	$1.091E-02$	$8.898E-03$

DOSE TO SELECTED ORGANS OF INTEREST VIA INHALATION
TASK GROUP ON LUNG DYNAMICS MODEL
DURATION OF INHALATION 6.00E+01 MIN
ORGAN DOSE IN REM

RADIO-NUCLIDE	DOSE TIME (DAYS)	KIDNEYS	LIVER	BONE	LUNGS
ALL	3.65E+02	4.08E-05	1.61E-04	2.01E-04	1.79E-02
ALL	1.83E+04	1.10E-02	3.59E-02	5.51E-02	4.50E-02

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