

Paper

RADIOACTIVITY COMPUTATION OF STEADY-STATE AND PULSED FUSION REACTORS OPERATION*

by

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June 1994

Submitted to the Third International Symposium on Fusion Nuclear Technology, University of California at Los Angeles, June 27 - July 1, 1994.

* Work supported by the United States Department of Energy/Office of Fusion Energy, under Contract No. W-31-109-Eng-38.

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Radioactivity Computation of Steady-State and Pulsed Fusion Reactors Operation *

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Abstract

Different mathematical methods are used to calculate the nuclear transmutation in steady-state and pulsed neutron irradiation. These methods are the Schür decomposition, the eigenvector decomposition, and the Padé approximation of the matrix exponential function. In the case of the linear decay chain approximation, a simple algorithm is used to evaluate the transition matrices.

1. Introduction

The International Thermonuclear Reactor (ITER) is expected to operate in a pulsed operational mode. Accurate radioactivity calculations, that take into account this mode of operation, are required in order to determine precisely the different safety aspects of ITER. We previously examined analytically the effect of pulsed operation in ITER and showed how it depends on the burn time, the dwell time, and the half-lives[0]. That analysis showed also that for ITER's low duty factor, using the continuous operation assumption would considerably overestimate the radioactivities, for a wide range of half-lives.

*Work supported by the U.S. Department of Energy, Office of Fusion Energy, under Contract No. W-31-109-Eng-38.

At the same time, the large improvements in the quality and the quantity (number of isotopes) of the decay and the cross-section data libraries[0] has considerably increased the computation times of the radioactivity calculations. For both reasons it is imperative to seek different methods of solution that reduce the computational time and can be easily adopted to the treatment of the pulsed operation.

In this work, we have developed algorithms based on several mathematical methods that were chosen based on their generality, reliability, stability, accuracy, and efficiency. These methods are the matrix Schür decomposition, the eigenvector decomposition, and the Padé approximation for the matrix exponential functions. In Section 2, the characteristics of the transmutation system of equation are discussed. Section 3 presents a brief review of the currently used methods. In Section 4, the new algorithms are presented.

2. The transmutation system of equations

The radioactivity inventory in a nuclear reactor is described by a set of simultaneous first-order linear differential equations. These equations can be written in a matrix notation as:

$$\dot{y} = Ay = (\phi C + D)y, \quad (2.1)$$

where

y = the concentration vector of all the nuclides involved,

ϕ = the position-averaged and energy-averaged neutron flux,

C = the microscopic cross-section matrix whose off-diagonal elements $c_{ij} = \sigma_{ij}$ are the spectrum-averaged microscopic cross-sections of the nuclide j forming nuclide i . The diagonal elements, $c_{ii} = -\sigma_i^t$, are the negative of the total destruction microscopic cross-section of the nuclide i ,

D = the decay matrix whose diagonal elements, d_{ii} , are the negative of the decay constant of the nuclides, and the off-diagonal elements, d_{ij} is the partial decay constant of the j^{th} nuclide decaying to the i^{th} nuclide,

$A = \phi C + D$ during operation, and $A = D$ during shutdown.

The elements of C and D are constants. The neutron flux may, however, change with time. In fission reactors the radioactive inventory is primarily due to the production of the fission products and thus depends strongly on the fuel burn-up that in turn changes the neutron flux. In fusion reactors, the neutron flux is due to the external plasma neutron source. As the materials interact with neutrons, the isotopes inventory changes. Such change could, in theory, alter the neutron spectrum. However, this change in the neutron spectrum, for all practical purposes, is small and can be neglected. As a result, the coefficients in Eq. 2.1 are considered constants.

The i^{th} row in Eq. 2.1 represents the rate of change of the i^{th} nuclide due to its destruction and the transmutations of the other nuclides to it. The off-diagonal elements of the j^{th} column in the matrix A are the different transmutations of the j^{th} nuclide to the other nuclides. Since a_{jj} is the negative of the total destruction cross-section, the summation of each column must equal zero, i.e. $\sum_i a_{ij} = 0$. Apart from the fission reaction, this condition implies the conservation of the total number of nuclides in the transmutation reactions. Accordingly, the sum of the concentration vector y at any time must equal the sum of the initial concentration vector, i.e. $\sum y_i(t) = \sum y_i(t_0)$. The decay matrix D is a triangular matrix, but the cross-section matrix C is, in general, not triangular, because of reactions such as $(n, 2n)$ that lead to feedback loops. Therefore, the diagonal elements of A are not generally identical to the eigenvalues of A .

The solution of Eq. 2.1 is given by [0]:

$$y(t) = \exp((t - t_0)A)y(t_0) = R(t, t_0)y(t_0), \quad (2.2)$$

where $y(t_0)$ is the initial concentration vector, and $R \equiv \exp(tA)$ is called the resolvent or the transition matrix. The function $\exp(tA)$ can be formally defined by the convergent power series:

$$\exp(tA) = I + tA + \frac{t^2 A^2}{2!} + \dots = \sum_{i=0}^{\infty} \frac{t^i A^i}{i!}, \quad (2.3)$$

where I is the identity matrix.

3. Methods of Solutions

The solution of Eq. 2.1 can be found with many different methods. For a review of many of these methods, the reader is referred to the article by Moler and Van

Loan[0]. In this section, we first review briefly historical and popular methods that have been used in the transmutation calculations. Then, we describe the matrix decomposition methods, the Padé approximation method, and a special method for lower triangular transmutation matrices.

3.1. Linear chain method

When Rutherford[0] established the possibility of the spontaneous transmutation of elements about a century ago, he deduced mathematically the solutions of the densities of the atoms involved in a successive radioactive-decay series. Rutherford considered limited cases in which there are only two products in addition to the primary substance. Bateman[0] introduced a more generalized mathematical method that can treat unlimited number of consecutive products.

The applicability of Bateman's method is confined to naturally occurring or artificially produced radioactive substances with linear successive transmutations ($A.B.C\dots$). The linearity, in this context, means a unique path from nuclide to nuclide with no feedbacks from daughters to parents. The linearity breaks down when the nuclides in the chain are subjected to nuclear reactions e.g. neutron reactions. In this case, different paths can lead to the same isotope with possible feedback (loops) from the products to their parents (e.g. $A(n, \gamma)B(n, 2n)A$; or $A(n, p)B(\beta^-)A$) or grandparents (e.g. $A(n, \gamma)B(n, d)C(\beta^-)A$). The system of differential equations then becomes highly coupled. This is indeed the case in fission and particularly in fusion reactors, where the high energy neutrons are capable of producing many loops.

In order to use the linear chain method to calculate the transmutations in a nuclear reactor, approximations have to be made. One may either neglect the loops altogether[0], or deal with only isolated first order loops[0], or ignore many nuclides and introduce fictitious chains and fictitious nuclides to decouple the differential equations[0]. These approximations can be justified for particular problems or chains, but can not be generalized.

3.2. Ordinary differential equation methods

Another approach for calculating the nuclear transmutation is the direct numerical integration of the radioactivity differential equations using an ordinary differential equation solver. The transmutation system of equations is an extremely stiff system with time constants that vary from fractions of a second to thousands of years. This requires the use of specialized methods such as the backward

differentiation formulas (BDF) used in the well-tested GEAR package[0] which has automatic step size control.

Such methods, however, are designed for general nonlinear differential equations and thus do not take advantage of the linearity and the constant coefficient nature of Eq. 2.1. Nonetheless, these methods provide more accurate accounting of the transmutations than the linear chain method. The radioactivity code RACC[0] uses the GEAR method. Another version of RACC[0] makes use of the LSODES solver[0] which is based also on the BDF method but designed for sparse systems.

3.3. Series Methods

The power series definition of $\exp(tA)$, Eq. 2.3, is the basis of these methods. Direct calculation of the series expansion is inefficient and unsatisfactory. Using the identity

$$e^A = (e^{A/m})^m, \quad (3.1)$$

can improve the efficiency and the accuracy of these methods[0]. This is done by scaling the matrix by m , which is taken as a power of two, $m = 2^k$, such that $\|A\|/m \leq 1$. The series is calculated for the scaled matrix and the result is then squared repeatedly. This scaling and squaring technique has been used in the ORIGEN code[0] to calculate Eq. 2.3.

3.4. Matrix decomposition methods

The matrix decomposition methods are based on the transformation of a matrix to a form that can be easily used. Generally this takes the form:

$$A = SBS^{-1}. \quad (3.2)$$

Using the above power series definition of e^{tA} , we find:

$$e^{tA} = Se^{tB}S^{-1}. \quad (3.3)$$

The closer B is to diagonal form, with invertible and well-conditioned S , the easier is the evaluation of e^{tB} .

3.4.1. Eigenvector decomposition

If A is an $n \times n$ matrix that has n independent eigenvectors $[v_1 | \dots | v_n] = V$ with corresponding eigenvalues $\lambda_1, \dots, \lambda_n$, then S and B can be set to V , and Λ , respectively. Where $\Lambda = \text{diag}(\lambda_1, \dots, \lambda_n)$. Then:

$$e^{tA} = V \text{diag}(e^{t\lambda_1}, \dots, e^{t\lambda_n}) V^{-1}. \quad (3.4)$$

The difficulties in this approach are that A does not always have a full set of independent eigenvectors and V , even if it is invertible, can behave very badly[0]. Another problem that has been encountered is that A can sometimes have complex eigenvalues thus forcing the costly use of complex arithmetic in Eq. 3.4.

3.4.2. Schür decomposition

For any real matrix A , there exists an orthogonal matrix Q (the Schür vectors of A), and a block upper triangular matrix T (the Schür form of A) such that:

$$Q^T A Q = \begin{bmatrix} T_{11} & T_{12} & \dots & T_{1n} \\ 0 & T_{22} & \dots & T_{2n} \\ \vdots & \vdots & \ddots & \vdots \\ 0 & 0 & \dots & T_{nn} \end{bmatrix} \quad (3.5)$$

where each T_{ii} is either a 1-by-1 block matrix or 2-by-2 matrix having complex conjugate eigenvalues. This means that $A = Q T Q^T$, and substituting this equation in Eq. 2.3, we get

$$e^{tA} = Q e^{tT} Q^T. \quad (3.6)$$

This requires the evaluation of the exponential of a triangular matrix.

Using the properties of triangular matrices, Parlett[0] showed that for any analytic function Φ of a block upper triangular matrix T , $\Phi(T) \equiv F$, is a also block upper triangular matrix with the same block structure as T and commutes with T , i.e.

$$F T = T F. \quad (3.7)$$

The diagonal blocks of F can be obtained from $F_{ii} = \Phi(T_{ii})$ and the upper diagonal blocks $i < j$ may be computed one diagonal at a time, moving out from the main diagonal, from the recursive relation:

$$T_{ii} F_{ij} - F_{ij} T_{jj} = \sum_{k=0}^{j-i-1} (F_{i,i+k} T_{i+k,j} - T_{i,j-k} F_{j-k,j}), \quad i < j. \quad (3.8)$$

If the transmutation matrix A has only real eigenvalues, T and F will have only 1×1 blocks, i.e. simple triangular matrices, and the computation of F is straightforward. If A possesses some complex eigenvalues, the F_{ij} blocks may be 1×1 , 2×1 , 1×2 , or 2×2 , depending on the structure of T . The 2×2 case requires the solution of four linear algebraic equations. The case of confluent eigenvalues requires special treatment[0] that uses the derivatives of the function Φ .

3.4.3. Matrix decomposition algorithm

The algorithm uses different routines from the mathematical library LAPACK[0]. The transmutation matrix is balanced and reduced to the Hessenberg matrix H , which is zero below the first subdiagonal, giving $A = GHG^T$, where G is orthogonal. The Hessenberg matrix is then reduced to the Schür form T as $H = STS^T$, where S is the Schür vectors of H . Thus, $A = GST(GS)^T$ and GS equals Q in Eq. 3.5.

At this point, all the eigenvalues of A are known (the diagonal of T). If required and if the eigenvalues are all real, the eigenvectors, may be computed and Eq. 3.4 is used to compute the resolvent. Otherwise, Eqs. 3.8 and 3.6 are used. In either case, most of the computation time is spent in the decomposition stage and is independent of the time t . Only at the last stage, the time t is used to compute e^{tT} (or e^{tA}). This makes these methods more valuable and amenable to the treatments of pulsed operation in terms of computational time saving as is discussed shortly.

3.5. The Padé approximation

In this work, the Padé approximation of the matrix exponential may be used to compute the exponential of the 2×2 diagonal blocks in Eq. 3.5, or may be used to calculate the resolvent e^{tA} directly. In both cases, the method proved to be efficient and reliable. The p^{th} diagonal Padé approximation of e^A is giving by:

$$N_p^{-1}(-A) N_p(A)$$

where

$$N_p(A) = \sum_{k=0}^p c_k A^k, \quad c_k = \frac{(2p-k)!p!}{(2p)!k!(p-k)!}$$

Ward[0] developed an algorithm that balances the matrix and makes use of the scaling and squaring technique (Eq. 3.1) to compute the exponential of an

arbitrary matrix using the above equation. In addition, the algorithm provides an estimate of the accuracy of the results. We have used Ward's algorithm and a modified version written by Van Loan[0] that provides more control on the degree of the approximation (p) and the accuracy.

3.6. Triangular systems

Naturally, if the transmutation matrix has a special shape, one can exploit such shape in order to reduce the computation time. As described above, during shut-down and in the linear chain method, the transmutation matrices are triangular having their eigenvalues on the diagonal. One can then make use of Eq. 3.8 directly to evaluate the resolvents. In fact, this equation is an elegant form of the Bateman equations that allows for the simultaneous solution of the system of equations, even with multiple paths to the same nuclide, without the need of the artificial break of the system as in the linear chain method.

Rewriting Eq. 3.8 for a simple lower triangular matrix, we have:

$$f_{ij} = \frac{1}{t_{ii} - t_{jj}} [t_{ij}(f_{ii} - f_{jj}) + \sum_{k=j+1}^{i-1} (f_{ik}t_{kj} - t_{ik}f_{kj})], \quad i > j. \quad (3.9)$$

Consider for example the following transmutation matrix and the corresponding resolvent

$$T = \begin{matrix} & \begin{matrix} y_1 & y_2 & y_3 \end{matrix} \\ \begin{matrix} y_1 \\ y_2 \\ y_3 \end{matrix} & \begin{pmatrix} -d_1 & 0 & 0 \\ d_{12} & -d_2 & 0 \\ d_{13} & d_2 & -d_3 \end{pmatrix} \end{matrix}, \quad F = e^{tT} = \begin{matrix} & \begin{matrix} y_1 & y_2 & y_3 \end{matrix} \\ \begin{matrix} y_1 \\ y_2 \\ y_3 \end{matrix} & \begin{pmatrix} f_{11} & 0 & 0 \\ f_{21} & f_{22} & 0 \\ f_{31} & f_{32} & f_{33} \end{pmatrix} \end{matrix}$$

truncated for only three isotopes. The diagonal elements f_{ii} are $f_{11} = e^{-td_1}$, $f_{22} = e^{-td_2}$, and $f_{33} = e^{-td_3}$. The off-diagonal elements can then be found recursively, one subdiagonal after another, using 3.9 as:

$$\begin{aligned} f_{21} &= \frac{t_{21}}{t_{22} - t_{11}} (f_{22} - f_{11}) = \frac{d_{12}}{d_1 - d_2} (e^{-td_2} - e^{-td_1}), \\ f_{32} &= \frac{t_{32}}{t_{33} - t_{22}} (f_{33} - f_{22}) = \frac{d_2}{d_2 - d_3} (e^{-td_3} - e^{-td_2}), \\ f_{31} &= \frac{t_{31}}{t_{33} - t_{11}} [t_{31}(f_{33} - f_{11}) + f_{32}t_{21} - t_{32}f_{21}] \\ &= \frac{1}{d_1 - d_3} [d_{13}(e^{-td_3} - e^{-td_1}) + \frac{d_{12}d_2}{d_2 - d_3} (e^{-td_3} - e^{-td_2}) - \frac{d_2d_{12}}{d_1 - d_2} (e^{-td_2} - e^{-td_1})]. \end{aligned}$$

The same procedure can be used to evaluate the power of the triangular matrix, say R^m , with the diagonal elements $f_{ii} = r_{ii}^m$.

4. Pulsed operation

For a few number of pulses, the nuclear transmutations can be computed by any of the methods described above by solving the transmutation equations for each pulse and shutdown period. For a large number of periodic sequences of pulses, one can save a significant amount of computing time by taking advantage of the regularity in the pulse sequence. This can be done if the solutions are obtained in terms of the resolvents of the different periods. As seen in Eq. 2.2, the resolvent is constant for the same transmutation matrix and the same time difference.

For a uniform sequence of pulses, suppose that $R_p(t_p)$ is the resolvent during the pulse time t_p , and $R_d(t_d)$ is the resolvent during the decay time t_d between pulses. Using Eq. 2.2 the solution at the end of t_d is

$$y(t_d + t_p) = R_d R_p y(0) \equiv R_c y(0), \quad (4.1)$$

where R_c is the resolvent for one cycle of a pulse and a decay period. If this cycle is repeated m times, the solution at the end of the m^{th} pulse is

$$y(mt_p + (m - 1)t_d) = R_p R_c^{m-1} y(0). \quad (4.2)$$

This requires at most $2[\log_2(m)]$ matrix multiplications[0].

Depending on the size of the matrix n and the number of pulses m , computing cost could be reduced by decomposing R_c itself using the matrix decomposition methods discussed above. Using the eigenvector decomposition

$$R_c^{m-1} = V \text{diag}(\lambda_1^{m-1} \dots \lambda_n^{m-1}) V^{-1} \quad (4.3)$$

where V and $\lambda_1 \dots \lambda_n$ are the eigenvectors and the eigenvalues of R_c , and using the Schür decomposition

$$R_c^{m-1} = Q T^{m-1} Q^T \quad (4.4)$$

where Q and T are the Schür vectors and form of R_c . The same matrix decomposition algorithm described above is used with the function $\Phi(x) = x^m$.

For triangular transmutation systems, R_p and R_d , are triangular and Eq. 3.9 can then be used to evaluate the powers of the resolvent of the cycle R_c , which is also triangular. Thus, the decomposition stage including the calculation of the eigenvalues of the triangular matrices[0] is unnecessary.

5. An example problem

The above algorithms described above have been coded for the CRAY computers which have 64-bit word and 47-bit mantissa. We demonstrate the results of the different methods for a problem used to compare different transmutation codes[0] utilizing reduced cross-section and decay data libraries and a specified neutron flux. The problem is the evaluation of the transmutation of ^{50}Cr after one year of irradiation. The initial atomic concentration of ^{50}Cr is set to 10. The transmutation matrix, with the reaction modes, is given by:

$$10^{-13} \begin{pmatrix} {}^{50}\text{Cr} & {}^{51}\text{Cr}^* & {}^{52}\text{Cr} & {}^{53}\text{Cr} & {}^{54}\text{Cr} & {}^{49}\text{V}^* & {}^{50}\text{V} & {}^{51}\text{V} & {}^{48}\text{T}_i & {}^{49}\text{T}_i \\ \begin{pmatrix} -2921 \\ \sigma_1 \\ 495.9 \\ n,\gamma \\ 914.8 \\ n,\gamma \\ -649.6 \\ \sigma_3 \\ 191.8 \\ n,\gamma \\ -548.3 \\ \sigma_4 \\ 546 \\ n,\gamma \\ 0 \\ 1197 \\ n,n,p \\ 1227 \\ n,p \\ 138.5 \\ n,n,p \\ 2897000 \\ c+n,p \\ 307 \\ n,\alpha \\ .006628 \\ n,^3\text{He} \\ .1178 \\ n,i \\ 314 \\ n,d \\ 2.321 \\ n,t \\ 429.4 \\ n,n,p \\ 243700 \\ \epsilon \\ 2442 \\ n,2n \\ -8846 \\ \sigma_7 \\ 5525 \\ n,7 \\ .1161 \\ n,n,d \\ 197.3 \\ n,np \\ 2.917 \\ n,i \\ 2667 \\ n,2n \\ -2667 \\ \sigma_{10} \end{pmatrix} \end{pmatrix}$$

Notice that the submatrix of the Cr isotopes is triangular and the analytical solutions of these isotopes can be found using Eq. 3.9. Table 5.1 shows this analytical solution, the results of the different methods, the deviations from the initial concentration giving in particles per million, and the CPU times in milli seconds. The results of the Cr isotopes from the different methods are shown as the percentage differences from the analytical solutions. For the other isotopes, the results are giving in absolute values.

For the Cr isotopes, the solutions of the matrix decomposition methods are the closest to the analytical solutions. The differences from the analytical solutions of these methods are orders of magnitude less than the other methods. The Schür method is the most accurate method. There is considerable reduction in the CPU times by using the decomposition methods and the Padé method. The LSODES method has the largest CPU time. However, this could be balanced, to some extent, by the fact that this method requires the minimum memory. It should

be noted that the CPU time depends on the size of the matrix and the saving in computing time could be greater for larger problems by using the decomposition methods.

Consider now the same problem for pulsed operation. To reach the same fluence used in the continuous irradiation calculations above, we assume 10000 pulses with a pulse width and a dwell time of 3156 and 2000 seconds, respectively. Table 2 shows the results, using the eigenvector decomposition, at the end of the first pulse, at the end of the first decay period, and at the end of the 10000th pulse. The computing time was only 31 ms, which is remarkably orders of magnitude less than that for a BDF method, which can be estimated from Table 1 as at least $10000 \times 2 \times 13 = 260000$ ms.

6. Conclusions

We have demonstrated the use of different mathematical methods for solving the transmutation system of equations in steady-state and pulsed operation fusion reactors. Considerable accuracy and efficiency can be achieved by using the matrix decomposition methods. A simple recursive relation has been developed to calculate the resolvents in triangular transmutation systems (linear chain methods). The results for a simple problem show the accuracy and the efficiency of the different methods, and the saving in computing times by using the matrix decomposition methods in pulsed operation.

Acknowledgments

The author wishes to acknowledge helpful and fruitful discussions with Dr. X. Sun of the Mathematical and Computer Science Division at Argonne National Laboratory, and Dr. J. Brooks of the Fusion Power Program at ANL for reviewing this paper.

Table 5.1: Solution Comparison- Transmutation of ^{50}Cr after one year of continuous operation^a

Isotope	Analytical	Schür	Eigenvector
^{50}Cr	9.90824823398	0.0 %	0.0 %
$^{51}\text{Cr}^*$	1.69666077856E-03	-1.77085549354E-10 %	-1.77085549354E-10 %
^{52}Cr	4.37693012077E-06	-1.13945844595E-10 %	-1.13945844595E-10 %
^{53}Cr	1.19872826563E-09	4.17120502630E-09 %	-2.26072577552E-07 %
^{54}Cr	6.29810678824E-13	-7.19772138009E-07 %	8.83054064462E-04 %
$^{49}\text{V}^*$	0.0	2.63103404748E-02	2.63103404748E-02
^{50}V	0.0	3.80864589509E-02	3.80864589509E-02
^{51}V	0.0	1.41676438411E-02	1.41676438410E-02
^{48}Ti	0.0	5.54928385814E-05	5.54928383281E-05
^{49}Ti	0.0	1.13900571692E-02	1.13900571714E-02
Error-ppm ^b	0.0	-4.07338371247	-4.07338350783
CPU(ms)	0	4	5
Isotope	Padé	GEAR	LSODES
^{50}Cr	0.0 %	0.0 %	0.0 %
$^{51}\text{Cr}^*$	6.30849143866E-05 %	-5.19315949417E-05 %	-4.94088313428E-07 %
^{52}Cr	-7.71967986919E-06 %	6.35502480098E-06 %	5.99739090522E-08 %
^{53}Cr	1.01357411537E-06 %	-2.39086670990E-06 %	-7.43287647805E-07 %
^{54}Cr	1.59468124902E-03 %	1.59707117337E-03 %	1.59571711576E-03 %
$^{49}\text{V}^*$	2.63103404748E-02	2.63103404719E-02	2.63103404678E-02
^{50}V	3.80864589516E-02	3.80864589504E-02	3.80864589502E-02
^{51}V	1.41676427705E-02	1.41676447224E-02	1.41676438497E-02
^{48}Ti	5.54928394367E-05	5.54928396125E-05	5.54928396451E-05
^{49}Ti	1.13900571682E-02	1.13900571712E-02	1.13900571751E-02
Error-ppm ^b	-4.07338370678	-4.07338372383	-4.07338367268
CPU(ms)	3	13	55

(a) Values in % are $(y_{\text{meth}} - y_{\text{anal}})/y_{\text{anal}} * 100$.

(b) Error= $10^8(\sum y_i(t) - \sum y_i(0))/\sum y_i(0)$.

Table 5.2: Transmutation of ^{50}Cr after 10000 pulses^a.

Isotope	End Pulse	End Dwell	10000 Pulse
^{50}Cr	9.999991	9.999991	9.908248
$^{51}\text{Cr}^*$	1.564205E-06	1.563300E-06	1.038884E-03
^{52}Cr	2.258176E-13	2.258176E-13	2.806988E-06
^{53}Cr	2.032879E-17	2.032879E-17	7.977416E-10
^{54}Cr	0.0	0.0	1.819256E-13
$^{49}\text{V}^*$	3.778731E-06	3.778547E-06	2.148223E-02
^{50}V	3.873765E-06	3.873765E-06	3.808704E-02
^{51}V	7.185008E-10	1.624175E-09	1.482433E-02
^{48}Ti	0.0	0.0	6.830789E-05
^{49}Ti	1.477929E-10	3.315157E-10	1.620743E-02
Error-ppm ^b	4.445155E-6	4.439471E-6	-4.072662
CPU(ms)	4	3	31

(a) Burn time=3156 seconds and dwell time=2000 deconds.

(b) Error= $10^6(\sum y_i(t) - \sum y_i(0))/\sum y_i(0)$.

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