Ptolemy
A Program for Heavy-Ion Direct-reaction Calculations
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Contents

1 Introduction 2

2 Summary of Formulas 5
  2.1 Two-Body Channels ................................................. 5
    2.1.1 Bound States ................................................. 5
    2.1.2 Scattering States ............................................ 6
  2.2 Inelastic Excitation .............................................. 8
    2.2.1 Differential Cross Section .................................. 9
    2.2.2 Geometrical Component and Effective Interactions ........... 10
    2.2.3 Strength of Effective Interaction ........................... 11
  2.3 Transfer Reactions ................................................ 11
    2.3.1 Possible Reactions ........................................... 12
    2.3.2 Two-Body States ............................................... 12
    2.3.3 Vector Transformation between Bound-State and Scattering Variables 15
    2.3.4 Calculation of Bound-State Form Factor ....................... 16
    2.3.5 DWBA Amplitude and Effective Interaction .................... 18
    2.3.6 Angular Momenta ............................................... 19
    2.3.7 Differential Cross Section ................................... 19
    2.3.8 Geometrical Component (including radial integrals) .......... 20
    2.3.9 Spectroscopic Factors and Spectroscopic Component .......... 20
    2.3.10 Outline of Steps in a DWBA Computation for Transfer Reactions 21
  2.4 Constants and Units ................................................ 22

3 Notation and Syntax 24
  3.1 Notation ............................................................. 24
  3.2 Ptolemy Syntax .................................................... 25

4 Elastic and Bound State Calculations 26
  4.1 Specifying the Two-Body Channel ................................ 26
  4.2 Specifying the Potentials ......................................... 28
  4.3 Controlling the Calculation ....................................... 31
  4.4 The Two-Body Wavefunctions ...................................... 33
  4.5 Reading the Output ................................................. 34
5 Optical Model Potential Fits
   5.1 Specifying the Fit Parameters and Data
   5.2 Controlling the Fit
   5.3 Reading the Output

6 Collective Model DWBA for Inelastic Excitation
   6.1 Specifying the Physical Problem
   6.2 Controlling the Calculation
   6.3 Reading the Output

7 Finite Range DWBA for Transfer Reactions
   7.1 Specifying the Physical Problem
   7.2 The Integration Grid
   7.3 Controlling the Calculation
   7.4 Saving Time and Core with SAVEHS and USEHS
   7.5 Reading the Output

8 Control Keywords

9 Acknowledgments

A List of all Keywords
   A.1 Reaction-defining Keywords
   A.2 Two-body State Keywords
   A.3 Calculation Keywords
   A.4 Control and Option-selecting Keywords
   A.5 Sub-keywords for DATA

B Sample Ptolemy Jobs
   B.1 Optical Model Fit
   B.2 Transfer Reaction
Abstract

PTOLEMY is a program for the computation of nuclear elastic and direct-reaction cross sections. It carries out optical-model fits to elastic-scattering data at one or more energies and for one or more combinations of projectile and target, collective model DWBA calculations of excitation processes, and finite-range DWBA calculations of nucleon-transfer reactions. It is fast and does not require large amounts of memory. The input is exceptionally flexible and easy to use. This report outlines the types of calculations that PTOLEMY can carry out, summarizes the formulas used, and gives a detailed description of its input.
Chapter 1

Introduction

PTOLEMY is a program for fitting optical-model potentials to elastic scattering data, and for the computation of the Distorted Wave Born Approximation to nuclear direct-reaction amplitudes. Either the collective-model DWBA for inelastic excitation or the finite-range DWBA for transfer reactions may be computed. No use is made of approximations that rely on the short range of nuclear interactions (e.g. zero-range and no-recoil approximations). PTOLEMY is specifically designed for heavy ion reactions but is nonetheless very efficient for light ion reactions. Advantages of PTOLEMY over other DWBA codes include high speed, low core requirements, and ease of use.

PTOLEMY derives its speed and compactness from several design features:

1. Substantial effort has been put into the development of the subroutine that picks the three-dimensional integration grid that is used in the DWBA transfer calculations. This subroutine makes use of the bound state form factor and the properties of the scattering wavefunctions. This efficiently chosen integration grid results in the need for relatively small numbers of integration points; as an example, a grid consisting of $24 \times 10 \times 10$ points will give accuracies of 1 or 2 percent for many heavy-ion reactions at moderate energies.

2. In high-energy heavy-ion DWBA transfer calculations, the scattering waves oscillate rapidly while the form factor varies slowly. However the computation of the form factor is the most time consuming aspect of the calculation. Therefore the form factor is computed on a coarse grid and interpolated to the finer grid needed for the integrals involving the scattering waves. This interpolation results in a reduction in the total computer time by a factor of two to five in typical heavy-ion calculations.

3. Interpolation and extrapolation in L-space is used to reduce the number of radial integrals that must be computed. Interpolation is achieved by fitting a continued fraction to the computed values, while the exponential form of the radial integrals for large $L$ is used for extrapolation. For Oxygen on Lead reactions, a time savings of 80% is realized by this method.

4. The calculation of the angular transforms in transfer reactions has been implemented using cosines instead of spherical harmonics. This approach avoids the
large numerical cancellations that occur in the more conventional Legendre decomposition methods if the exchanged orbital angular momentum is relatively large. A specially designed in-line cosine routine and iterative evaluations of cosines are used to reduce the cosine computation time.

5. The Coulomb terms of the inelastic excitation amplitudes are computed using recursion relations in $L$. The starting values for the recursion relations are generated using Belling’s asymptotic expansion for integrals of Coulomb wavefunctions and powers of $r$. A new asymptotic expansion of Coulomb wavefunctions for large argument is used in these calculations. The result of these techniques is a high speed program for the Coulomb excitation in which it is not necessary to be concerned with either orbital angular momentum or radial cut-offs.

6. The optical-model potential fitting part of PTOLEMY uses state-of-the-art minimization routines that make specific use of the sum of squares property of the function ($\chi^2$) being minimized. The gradients required by these minimizers are computed analytically.

7. The computation of the transfer amplitudes has been factored into segments in such a way that the recomputation of the same quantities is held to a reasonable (although not absolute) minimum without the need for extremely large tables. Scratch files are not used in any part of the program.

8. All arrays used in a PTOLEMY calculation are stored in a section of memory referred to as the allocator. Space in the allocator is reused when the data contained in it is no longer needed.

The result of these and other features is a very fast program for the finite-range DWBA. Calculations require only a matter of seconds on modern computers.

In addition to its great speed, PTOLEMY provides the user with an especially simple form of input. The input is designed to be flexible, tolerant of minor syntactical variations, concise and easy to remember. Extensive checking of the input is carried out in an effort to allow the same problem to be stated in a variety of ways and to eliminate the chance of calculations being made with undefined or otherwise unexpected parameters. In the following sections it will become evident that many quantities may be defined in more than one way. Often there will be the possibility of a direct specification of a parameter that can also be determined by PTOLEMY from other input. In most cases if the user explicitly specifies the value, PTOLEMY will not use the indirectly specified value nor will it check the two values for consistency. Therefore one should avoid needless duplication of input so that inadvertent inconsistencies do not occur. An example would be the specification of $Q$ after the bound state calculations – the new value of $Q$ would be used in determining the outgoing state scattering energy even though it might be inconsistent with the difference of the bound state energies. (In this case a warning message would be printed.)

Provision is made for keeping final results (radial integrals, elastic S-matrix elements, differential cross sections, etc.) in a form suitable for subsequent processing with
Speakeasy\textsuperscript{1} This is particularly useful for the production of graphs showing the results of one or more DWBA calculations. In addition one may use the extensive facilities of Speakeasy to manipulate cross sections or radial integrals interactively.

In addition to the computation of DWBA cross sections, PTOLEMY can also be used to fit optical potentials to elastic scattering data. The specification of the parameters to be varied is both simple and flexible; the user does not need to write a subroutine for each fit to be made. The method of entering data is quite general; if the user's data is already in a computer file, he or she will probably not need to reenter it for PTOLEMY. Fits to data at several energies may be made, and several keywords are provided to give the optical potential a dependence on the scattering energy. The user is given a choice of six different minimum-search programs including four that use analytically computed gradients. Two of the latter fitters work exceptionally well and 12-parameter fits to $^{16}$O+$^{208}$Pb data at five different energies may be made in only seconds.

In the next section we present the formulas used in PTOLEMY. The syntax used in PTOLEMY's input is described in Chapter 3. Chapter 4 describes elastic-scattering and bound-state calculations and contains basic material (such as potential definitions) that is used in the subsequent sections. The next three chapters (5, 6, and 7) discuss optical model fits, inelastic excitation, and transfer reactions respectively. These three chapters are largely independent of each other; where necessary, cross references are made. Chapter 9 presents some of the control keywords that perform functions auxiliary to the calculational functions. Appendix A contains a complete list of PTOLEMY keywords and their default values. Appendix B contains the complete input decks for some sample jobs.

This manual does not enumerate all possible variants and interpretations of the PTOLEMY input; rather it is limited to the most straightforward methods of stating the problem to be solved. For many DWBA calculations and optical-model fits, the necessary input can be inferred directly from the examples given in Appendix B, to which the tyro is referred. The only aspect of the input that is not clear upon inspection of these examples is the use of the \texttt{PARAMETERSET} keyword which is discussed on Section 4.3.

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Chapter 2

Summary of Formulas

This chapter contains a summary of the formulas used in PTOLEMY for evaluating elastic, inelastic and transfer cross sections. Since all of the formulas are standard, no attempt is made to present derivations.

2.1 Two-Body Channels

We will consider two particles (nuclei) referred to as the “projectile” (subscript p) and the “target” (subscript t). For scattering states these designations will have their customary meanings, while for bound states the only distinction is that the spin of the projectile determines the spin-orbit force. The reduced mass is

$$M = \frac{M_p M_t}{M_p + M_t} M_{\mu},$$

(2.1)

where $M_p$ and $M_t$ are the atomic weights of the two nuclei and $M_{\mu}$ is the atomic mass unit [Eq. 2.100].

2.1.1 Bound States

The Schrödinger equation for bound states may be written as

$$\left[ -\frac{1}{r^2} \frac{d}{dr} r^2 \frac{d}{dr} + \frac{\ell(\ell + 1)}{r^2} + \frac{2M}{\hbar^2} V(r) + \kappa^2 \right] \phi(r) = 0,$$

(2.2)

where

$$\kappa = \sqrt{-2ME/\hbar},$$

(2.3)

and $E$ is the (negative) energy of the bound state. For large $r$, $\phi(r)$ tends to a Whittaker function:¹

$$\phi(r) \xrightarrow{r \to \infty} \frac{N}{r} W_{-\eta,\ell+1/2}(2\kappa r),$$

(2.4)

¹M.Abramovitz and I.A.Stegun, Handbook of Mathematical Functions (Dover, New York, 1965)
where \( N \) is a normalization constant and \( \eta \) is the Sommerfeld parameter for the bound state:

\[
\eta = Z_p Z_t \alpha M c/(\hbar \kappa). \tag{2.5}
\]

The relevant combinations of the fine-structure constant \( \alpha \), the speed of light \( c \), and the Planck constant \( \hbar \) are given in Eqs.2.99 – 2.101. The bound state wave function is real and normalized such that

\[
\int_0^\infty dr r^2 |\phi(r)|^2 = 1. \tag{2.6}
\]

The sign of \( \phi(r) \) follows the conventions of Meyer and Jensen, \(^2\) namely \( \phi(r) \) is positive for small \( r \).

### 2.1.2 Scattering States

The Schrödinger equation for scattering states is written as

\[
\left[ -\frac{d^2}{dr^2} + \frac{\ell (\ell + 1)}{r^2} + \frac{2M}{\hbar^2} V(r) - k^2 \right] f_\ell(r), \tag{2.7}
\]

where the wave number is

\[
k = \sqrt{2M E}/\hbar, \tag{2.8}
\]

with \( E \) the center-of-mass scattering energy. If the potential contains a spin-orbit component, the \( f_\ell \) should also contain a label for the total projectile angular momentum; we will for the moment suppress this label. The scattering wavefunctions are normalized such that

\[
f_\ell(r) \xrightarrow{r \to \infty} \frac{1}{2} \left[ (1 + S_\ell) F_\ell(\eta, kr) + i(1 - S_\ell) G_\ell(\eta, kr) \right] = e^{+\delta}(\cos \delta F + \sin \delta G), \tag{2.9}
\]

where \( F_\ell \) and \( G_\ell \) are the regular and irregular Coulomb wavefunctions \(^3\) and \( \eta \) is the Sommerfeld parameter:

\[
\eta = Z_p Z_t \alpha M c/(\hbar k), \tag{2.10}
\]

\[
= Z_p Z_t \alpha \sqrt{M c^2 / 2E}. \tag{2.11}
\]

The elastic scattering \( S \)-matrix element \( S_\ell \) is defined by Eq. 2.29.


2.1. TWO-BODY CHANNELS

If there are no spin-dependent forces and if the particles are not identical, the elastic scattering cross section is given by

\[ \frac{d\sigma}{d\Omega} = |F(\theta)|^2. \] (2.12)

If the particles are identical, but there are no spin-dependent forces, the cross section is

\[ \frac{d\sigma}{d\Omega} = \left| F^{\pm}(\theta) \right|^2 \mp \frac{S_p}{S_p + 1/2} \left[ \left| F^{+}(\theta) \right|^2 - \left| F^{-}(\theta) \right|^2 \right], \] (2.13)

where the upper signs are for Bose statistics and the lower signs are for Fermi statistics. Here \( S_p \) is the spin of the particles. Finally, for a spin-1/2 projectile interacting with a spinless target via a spin-orbit force, the spin-averaged cross section is

\[ \frac{d\sigma}{d\Omega} = |F(\theta)|^2 + |B(\theta)|^2. \] (2.14)

We do not consider more complicated systems with spin-dependent forces.

The amplitudes in the above equations are sums of Rutherford and nuclear components,

\[ F(\theta) = F_R(\theta) + F_N(\theta), \] (2.15)

\[ F_R(\theta) = -\frac{\eta}{2k|\sin(1/2\theta)|^2} e^{2i[\sigma_0 - \eta \ln \sin(\frac{1}{2}\theta)]}, \] (2.16)

\[ F_N(\theta) = \frac{1}{2ik(2S_p + 1)} \sum_{j} (2j + 1)(S_{\ell,j} - 1)e^{2i\sigma_\ell} P_{\ell,0}(\cos \theta), \] (2.17)

\[ F^{\pm}(\theta) = F(\theta) \pm F(\pi - \theta), \] (2.18)

and

\[ B(\theta) = \frac{1}{2ik} \sum_{\ell} [S_{\ell,\ell+1/2} - S_{\ell,\ell-1/2}] e^{2i\sigma_\ell} P_{\ell,1}(\cos \theta). \] (2.19)

Here \( S_p \) is the spin of the projectile and the S-matrix elements have been labeled with both \( \ell \) and \( j \) to allow for a spin-orbit interaction. If there is no spin-orbit force, one has

\[ \frac{1}{2S_p + 1} \sum_{j} (2j + 1)S_{\ell,j} = (2\ell + 1)S_\ell. \] (2.20)

The Coulomb phase shifts are

\[ \sigma_\ell = \arg \Gamma(\ell + 1 + i\eta), \] (2.21)

and the conventions of Abramovitz and Stegun are used for the Legendre functions \( P_{\ell,m} \).

In the Rutherford amplitude (Eq. 2.16) an alternative form is given by the replacement

\[ \frac{\eta}{2k} = Z_p Z_t \alpha hc/(4E), \] (2.22)
The total reaction cross sections are given by
\[ \sigma_{\text{Reac}} = \frac{\pi}{k^2(2S_p + 1)} \sum_{\ell,j} (2j + 1)(1 - |S_{\ell,j}|^2) \] (2.23)
for non-identical particles and by
\[ \sigma_{\text{Reac}} = \frac{2\pi}{k^2(2S_p + 1)} \left( S_p \sum_{\ell} (2\ell + 1)(1 - |S_{\ell}|^2) + \sum_{\ell \text{ even or odd}} (2\ell + 1)(1 - |S_{\ell}|^2) \right) \] (2.24)
for identical particles. Here the second sum is over even partial waves for Bose statistics and over odd partial waves for Fermi statistics.

One can define a “nuclear total cross section”
\[ \sigma_{\text{Nuc}} = \frac{4\pi}{k} \text{Im} F_N(0) \] (2.25)
for non-identical particles, and
\[ \sigma_{\text{Nuc}} = \frac{4\pi}{k(2S_p + 1)} \left[ 2S_p \text{Im} F_N(0) + \text{Im} F_N^\pm(0) \right] \] (2.26)
for identical particles. If there is no Coulomb force, these quantities are just the total cross sections:
\[ \sigma_{\text{Nuc}} = \sigma_{el} + \sigma_{\text{Reac}} \] (2.27)
where \( \sigma_{el} \) is the integral over angles of the elastic cross section. Schwarzchild et.al. have discussed the significance of \( \sigma_{\text{Nuc}} \) in the presence of a Coulomb force; for heavy-ion scattering \( \sigma_{\text{Nuc}} \) is usually quite small.

### 2.2 Inelastic Excitation

We consider the inelastic excitation process
\[ A(a, b)B \] (2.28)
with either
\[ b = a, B = A^*, \] (2.29)
for target excitation or
\[ b = a^*, B = A, \] (2.30)
for projectile excitation.

The outgoing kinetic energy in the c. m. system is
\[ E_{\text{out}} = E_i + Q \] (2.31)
2.2. **INELASTIC EXCITATION**

where $E_i$ is the incoming c. m. energy and the $Q$ value is given by

$$Q = E_A + E_a - E_B - E_b$$

(2.32)

where the $E_k$ are the total energies (in MeV) of the nuclear states involved:

$$E_k = (Z_k M_p + N_k M_n) c^2 + (M_{xc})_k + E^*_k$$

(2.33)

or

$$E_k = E_g(Z_k, N_k) + E^*_k$$

(2.34)

$Z_k, N_k$ are the numbers of nucleons in the nuclear state $k$; $(M_{xc})_k$ is the ground state mass excess of the nucleus with $Z_k$ protons and $N_k$ neutrons and $E^*_k$ is the excitation energy if state $k$ is excited. $E_g(Z_k, N_k)$ is the total ground-state energy (or mass $\times c^2$) of the nucleus $(Z_k, N_k)$.

**PTOLEMY** evaluates the inelastic cross sections in DWBA using collective-model form factors for the nuclear part of the excitation. The distorted waves are the solution of Eq. 2.7 the normalization of Eq. 2.9. These distorted waves must, of course, be separately evaluated in the incoming $[a+A]$ and outgoing $[b+B]$ channels.

### 2.2.1 Differential Cross Section

The differential cross section at a c.m. scattering angle $\theta$ for the excitation process is written as

$$\frac{d\sigma}{d\Omega} = \frac{1}{E_i E_{out}} \frac{k_{out}}{k_i} R \sum_{L_x M_x} \left| G_{L_x M_x}^{L_x}(\theta) \right|^2$$

(2.35)

where $R$ is a spin-statistical factor:

$$R = \begin{cases} \frac{2J_b + 1}{2J_a + 1} & \text{(projectile excitation)} \\ \frac{2J_B + 1}{2J_A + 1} & \text{(target excitation)} \end{cases}$$

(2.36)

Here $k_i$ and $k_{out}$ are the wave numbers (Eq. 2.8 in the incoming and outgoing channels and $J_a, J_b, J_A,$ and $J_B$ are the intrinsic angular momenta of the four nuclei. The multipolarity of the transition is labeled by $L_X$.

The multipole transition amplitude may be decomposed into a term reflecting the amount of deformation and a geometric term:

$$G_{M_x L_x}^{L_x}(\theta) = A_{L_x} B_{M_x L_x}^{L_x}(\theta)$$

(2.37)
2.2.2 Geometrical Component and Effective Interactions

The “geometrical” component is

\[ B_{Mx}^{L_z} = \sqrt{4\pi} i^{L_z} \sum_{L_i L_{out}} C_{M_x,-M_x,0}^{L_{out},L_z,L_i} \times I_{L_i,L_{out},L_x} Y_{L_{out}}^{L_x}(\theta,0) \]  

(2.38)

where \( Y_{M_x}^{L_z}(\theta,\phi) \) is a spherical harmonic and \( C_{M_1,M_2,M}^{L_1,L_2,L} \) is a Clebsch-Gordan coefficient, both defined according to Condon-Shortley conventions.

The radial integrals are given by

\[ I_{L_i,L_{out},L_x} = \sqrt{\frac{2L_{out}+1}{4\pi}} \left| C_{0,0,0}^{L_{out},L_x,L_i} \right| e^{i(\sigma_i + \sigma_{out})} \times \int_0^{\infty} f_{L_i}(r) H_{L_x}(r) f_{L_{out}}(r) dr \]  

(2.39)

The symbols \( \sigma_i \) and \( \sigma_{out} \) designate the Coulomb phase shifts \( \sigma_{L_i}(\eta_i) \) and \( \sigma_{L_{out}}(\eta_{out}) \) respectively. The effective interaction \( H \) contains both nuclear and electric Coulomb contributions:

\[ H_{L_x}(r) = H_{L_x,N}(r) + H_{L_x,C}(r) \]  

(2.40)

The nuclear component of the effective interaction is

\[ H_{L_x,N}(r) = \frac{\beta_{L_x}}{\beta_{L_x}} \left[ R' \frac{dV(r)}{dr} + iR'_I \frac{dV_I(r)}{dr} \right] \]  

(2.41)

where \( V \) and \( V_I \) are the real and imaginary parts of an optical potential. The radii \( R' \) and \( R'_I \) are the radii of the excited nucleus; specifically:

\[ R' = r_0 A''^{(1/3)} \]
\[ R'_I = r_{I0} A''^{(1/3)} \]  

(2.42)

where \( A'' = A \) for target excitation, \( A'' = a \) for projectile excitation. The nuclear deformation parameter is \( \beta_{L_x} \) so the \( R' \beta_{L_x} \) and \( R'_I \beta_{L_x} \) are the deformation lengths. We have normalized the effective interaction to the average of the nuclear and Coulomb deformation parameters:

\[ \bar{\beta}_{L_x} = 1/2(\beta_{L_x} + \beta_{L_x,C}) \]  

(2.43)

The Coulomb part of the effective interaction is derived from the multipole expansion of the potential between a point charge and uniformly charged sphere.

\[ H_{L_x,C}(r) = \frac{\beta_{L_x,C}(R'_C)^3}{\beta} \frac{3Z_a Z C e^2}{2L_x + 1} \begin{cases} \frac{R_{C}^{2 L_x + 1}}{r} & r < R_C \\ \frac{R'_I^{L_x}}{r^{L_x + 1}} & r > R_C \end{cases} \]  

(2.44)
2.3. TRANSFER REACTIONS

where \( Z_a \) and \( Z_A \) are the atomic numbers of the two nuclei. Note the distinction between the Coulomb radius of the excited nucleus:

\[
R'_C = r_{C0} A^{(1/3)}
\]  

(2.45)

and the Coulomb radius of the optical potential:

\[
R_C = r_{C0} (a^{1/3} + A^{1/3})
\]  

(2.46)

The Coulomb deformation parameter \( \beta_{L_x,C} \) is related to the reduced transition rate \( B(E, L_x, \uparrow) \) by

\[
B(E, L_x, \uparrow) = \frac{2J_{\text{final}} + 1}{2J_{\text{initial}} + 1} B(E, L_x, \downarrow)
\]

\[
= \left[ \frac{3Z R'_{L_x}}{4\pi 10} \beta_{L_x,C} \right]^2 \times \frac{(2J_{\text{final}} + 1)}{(2J_{\text{initial}} + 1)(2L_x + 1)}
\]

(2.47)

where \( B(E, L_x, \uparrow) \) is given in units of \( e^2 \text{barn}^{L_x} \) and \( R'_C \) is in fm. The \( \beta_{L_x,C} \) so defined can be related to a nuclear deformation only for \( 0^+ \) ground states; however PTOLEMY correctly computes the Coulomb excitation from other ground states if the \( B(E, L_x, \uparrow) \) is given.

2.2.3 Strength of Effective Interaction

The strength constant in Eq. 2.37 is given by

\[
A_{L_x} = \frac{\beta_{L_x}}{\sqrt{2L_x + 1}}
\]

(2.48)

where \( \beta_{L_x} \) is defined in Eq. 2.43.

2.3 Transfer Reactions

PTOLEMY computes amplitudes and cross sections of nucleon-transfer reactions

\[
A(a, b)B
\]

(2.49)

using the full distorted-wave Born approximation (DWBA), without further approximations based on the short range of nuclear reactions. In the incoming channel \((A, a)\), \( A \) is the target and \( a \) is the projectile; in the outgoing channel \((B, b)\), \( B \) is the residual nucleus and \( b \) is the ejectile. \( A, a, B, b \) will be used both as identifiers of the nuclear states involved and as symbols for the total number of nucleons. The group of nucleons transferred (or the number of transferred nucleons) will be denoted by \( x \).
2.3.1 Possible Reactions

(i) If $a > b$, the reaction (Eq. 2.49) is a stripping reaction shown in Fig. 2.1.

\[
a = b + x \quad B = A + x
\]

(ii) If $a < b$, the reaction (Eq. 2.49) is a pick-up reaction as shown in Fig. 2.2.

\[
b = a + x \quad A = B + x
\]

2.3.2 Two-Body States

Calculation of the transfer cross sections involves the combination of four elements – the scattering wave functions in incoming (\(i\)) and outgoing (\(out\)) channels and bound-state wave functions representing the composite nucleus at each reaction vertex. The vertex involving $A$, $B$, and $x$ will be referred to as the target vertex and the corresponding bound state as the target bound state; the vertex involving $a$, $b$, and $x$ will be referred to as the projectile vertex and the corresponding bound state the projectile bound state.

---

2.3. TRANSFER REACTIONS

The kinetic energies in the incoming and outgoing channels are related by Eq. 2.31 – 2.34 and the inelastic scattering distorted waves for these two channels are found using Eq. 2.27 – 2.29; the radial variable in these equations is the position vector of the mass center of \(a\) (or \(b\)) relative to the mass center of \(A\) (or \(B\)). We will also need elastic scattering wavefunctions in incoming and outgoing channels,

\[
\chi^+(\vec{k}_i, \vec{r}_i) = \frac{4\pi}{k_i r_i} \sum_{L_i} i^{L_i} f_{L_i}(r_i) \left[ Y^{L_i}(\hat{r}_i) \cdot Y^{L_i}(\hat{k}_i) \right] (2.52)
\]

and

\[
[\chi^- (\vec{k}_{out}, \vec{r}_{out})]^* = \frac{4\pi}{k_{out} r_{out}} \sum_{L_{out}} i^{L_{out}} f_{L_{out}}[Y^{L_{out}}(\hat{r}_{out}) \cdot Y^{L_{out}}(\hat{k}_{out})] (2.53)
\]

Consider next the interaction vertices.

**The Target Vertex:** \( t = (x, C_t) \)

![Figure 2.3: The target vertex](image)

Let \( C_t \) denote the core nucleus and \( H_t \) the composite (heavier) nucleus at the target vertex as shown in Fig. 2.3

\[
C_t = B \quad (\text{pick-up}) \quad H_t = A \quad (\text{pick-up})
\]

\[
= A \quad (\text{stripping}) \quad = B \quad (\text{stripping})
\]

In either case vertex \( t \) is regarded as the break-up of the bound state \( H_t \) into its constituents

\[
H_t \rightarrow C_t + x
\]

(2.54)

The radial variable associated with this break-up is

\[
\vec{r}_t = \vec{r}_{c,x} = \vec{r}_{Bx} \quad (\text{pick-up})
\]

\[
= \vec{r}_{Ax} \quad (\text{stripping})
\]

(2.55)

Introduce a complete set of states

\[
\Psi_{m_t}^{n_t, \ell_t} (\vec{r}_t) = \Phi_{m_t \ell_t} (r_t) Y_{m_t}^{\ell_t} (\hat{r}_t)
\]

(2.56)

describing the bound-state wave-function of \( x \) and \( C_t \). Here \( \phi_{m_t \ell_t} \) is the bound-state radial wavefunction defined in Eq. 2.22 – 2.24. Let the transferred nucleons have intrinsic spin.
The particles emitted (or absorbed) at vertex \( t \) are described in terms of the functions

\[
[\Psi^{n_t\ell_t}(\mathbf{r}_t) \times \Phi^{xJ_x}(\rho_x)]_{J_t}^{M_t}
\]

(2.57)

where \( \Phi^{xJ_x} \) represents the intrinsic structure of \( x \).

The break-up of the internal wave function of \( H_t \) into \( C_t + x \) is to be described in terms of shell-model wave functions. This is characterized by a spectroscopic amplitude \( \Theta(n_t\ell_t, xJ_x, J_t; H_t, C_t) \) defined below [Eq. 2.96 and 2.97]. The coupling schemes for the angular momenta at vertex \( t \) are

\[
\vec{\ell}_t + \vec{J}_x = \vec{J}_t
\]

(2.58)

\[
\vec{J}(t) + \vec{J}_t = \vec{J}(H_t)
\]

(2.59)

Equation 2.50 defines the total angular momentum \( J_t \) transferred at vertex \( t \); Eq. 2.56 specifies how \( J_T \) is divided into orbital and intrinsic components.

**Projectile Vertex:** \( p = (x, C_p) \)

\[
\begin{array}{c}
\text{H}_p \\
\hline
\text{C}_p
\end{array}
\]

\[
\xrightarrow{x}
\]

Figure 2.4: Projectile Vertex

As in the treatment of the target vertex, \( C_p \) denotes core, \( H_p \) composite (heavier) nucleus. The projectile vertex is shown in Fig. 2.4.

\[
C_p = \begin{array}{ll}
a & \text{(pick-up)} \\
b & \text{(stripping)}
\end{array} \\
H_p = \begin{array}{ll}
b & \text{(pick-up)} \\
a & \text{(stripping)}
\end{array}
\]

Thus the vertex \( p \) describes the break-up of the bound state \( H_p \) into its constituents

\[
H_p \rightarrow C_p + x
\]

(2.60)

The radial variable is

\[
\vec{r}_p = \vec{r}_{C_px} = \vec{r}_{ax} \quad \text{pick-up}
\]

\[
= \vec{r}_{bx} \quad \text{stripping}
\]

(2.61)

(2.62)

The equations describing the state of the particles emitted (or absorbed) at \( p \) are obvious analogs of Eq. 2.56 and 2.57. The equations analogous to Eq. 2.58 and 2.59 for the angular momentum \( \vec{J}_p \) transferred at vertex \( p \) are:

\[
\vec{\ell}_p + \vec{J}_x = \vec{J}_p
\]

(2.63)

\[
\vec{J}(C_p) + \vec{J}_p = \vec{J}(H_p)
\]

(2.64)

\[^5\]Also used for the number of transferred nucleons.
2.3. TRANSFER REACTIONS

2.3.3 Vector Transformation between Bound-State and Scattering Variables

Ignoring for the moment the internal structure of the nuclear states $C_p$ and $C_t$, the transfer reactions under consideration are $(x + 2)$-body processes – the "bodies" are the two cores and the $x$ transferred nucleons. Now let the effective interaction that induces transfer be taken to be a function of the bound-state variables $\vec{r}_p$, $\vec{r}_t$ only; i.e., it is independent of the internal coordinates $\rho_x$ of $x$, dependent only on the position of the mass center of the $x$ transferred nucleons. The $(x + 2)$-body process now becomes a 3-body process $(C_p, C_t, x)$.

The natural variables for this 3-body problem are the position vectors of $C_p$, $C_t$, and $x$ relative to an origin in fixed space. In order to separate the center-of-mass motion, introduce as independent variables the center-of-mass position vector $\vec{R}$ and two (any two) of the relative variables $\vec{r}_i$, $\vec{r}_{out}$, $\vec{r}_t$, $\vec{r}_p$. PTOLEMY uses the scattering variables ($\vec{r}_i$, $\vec{r}_{out}$) as integration variables. The Jacobian of the transformation

$$(\vec{R}_{Ct}, \vec{R}_{Cp}, \vec{R}_x) \rightarrow (\vec{R}, \vec{r}_i, \vec{r}_{out})$$

is

$$J = \alpha^3$$

where

$$\alpha = \frac{H_p H_t}{(C_p + C_t + x)x}$$

(2.66)

and is included in the expression Eq. 2.67 for the transformation function $H$.

With $\vec{r}_i$, $\vec{r}_{out}$ as independent variables, angular-momentum functions of $\vec{r}_p$, $\vec{r}_t$ must be expressed in terms of $\vec{r}_i$, $\vec{r}_{out}$. This involves evaluation of the coefficients $H_{L_i,L_{out},L_x}^{n_{it},n_{tp},\ell_{tp}}(r_i, r_{out})$ of the vector transformation

$$\alpha^3 \Phi_{n_{it},\ell_{ti}}(r_t) V_{eff} \Phi_{n_{tp},\ell_{tp}}(r_p) \left[ Y^{\ell_{ti}}(\hat{r}_t) \times Y^{\ell_{tp}}(\hat{r}_p) \right]_{L_x}^{Lz}$$

$$= \sum_{L_i, L_{out}} (-)^{1/2(L+L_{out}+\ell_p-\ell_t)} H_{L_i,L_{out},L_x}^{n_{it},n_{tp},\ell_{tp}}(r_i, r_{out}) \left[ Y^{L_i}(\hat{r}_i) \times Y^{L_{out}}(\hat{r}_{out}) \right]_{M_x}^{Lz}$$

(2.67)

of the bound-state product into spherical harmonics of the independent variables $\vec{r}_i$, $\vec{r}_{out}$. Here $V_{eff}$ is the effective interaction that induces the transition. It will be defined below [Eq. 2.85 and 2.86]; all that is of consequence here is that $V_{eff}$ depends only on the radial variables. $H(r_i, r_{out})$ is referred to as the bound-state form factor. The strange phase factor in Eq. 2.67 is introduced to give the phase of the radial integrals $I_{L_i,L_{out},L_x}$ [Eq. 2.93] the following two desirable features for large values of $L_i$ and $L_{out}$ (suppressing for the moment the factor $e^{i(\sigma_i + \sigma_{out})}$):

1. The phase does not change by $\pi$ every time $L_i$ and $L_{out}$ change by unity.

2. For most reactions, the phase tends to 0 from above as both $L_i$ and $L_{out}$ increase.
2.3.4 Calculation of Bound-State Form Factor

The procedure for computation of the form factor $H$ in (Eq. 2.67) is that of Balian and Brézin. First, $\vec{r}_t$ and $\vec{r}_p$ must be expressed in terms of $\vec{r}_i$ and $\vec{r}_{out}$

\[
\begin{pmatrix}
\vec{r}_t \\
\vec{r}_p
\end{pmatrix}
= \begin{bmatrix}
s_1 & t_1 \\
s_2 & t_2
\end{bmatrix}
\begin{pmatrix}
\vec{r}_i \\
\vec{r}_{out}
\end{pmatrix}
\tag{2.68}
\]

To identify the constants $s_i, t_i$ consider the plane triangle whose vertices are the three basic particles $C_t, C_p, x$.

Figure 2.5: Coordinates for bound-state form factor

$H_t$ and $H_p$ are at the mass centers of $(C_p, x)$ and $(C_p, x)$. The vectors $\vec{r}_t$ and $\vec{r}_{out}$ have different identification for pick-up [Fig. 2.1] and stripping [Fig. 2.2]:

\[
\begin{align*}
\vec{r}_i &= \vec{r}_{Aa} = \vec{r}_{HtC_p} \\
\vec{r}_{out} &= \vec{r}_{Bb} = \vec{r}_{C_pH_p} \quad \text{pickup} \tag{2.69} \\
\vec{r}_i &= \vec{r}_{Aa} = \vec{r}_{CtH_p} \\
\vec{r}_{out} &= \vec{r}_{Bb} = \vec{r}_{HtC_p} \quad \text{stripping} \tag{2.70}
\end{align*}
\]

Then with $\alpha$ defined by Eq. 2.66 and

\[
\gamma = \frac{C_p}{H_p} \quad \delta = \frac{C_t}{H_t} \tag{2.71}
\]

\[
\begin{bmatrix}
s_1 \\
\delta
\end{bmatrix}
= \alpha
\begin{bmatrix}
-\gamma & 1 \\
-1 & \delta
\end{bmatrix}
\begin{cases}
\text{pick - up} \\
\text{stripping}
\end{cases}
\]

\[
\begin{cases}
\begin{align*}
\alpha &= \frac{1}{(a + B + x)x} \\
\alpha &= \frac{1}{(A + b + x)}
\end{align*} \tag{2.72}
\end{cases}
\]
2.3. TRANSFER REACTIONS

The bound-state form factor is then given by

\[
H_{L_iL_{out}L_x}^{n\ell_t\ell_p\mu_p}(r_i,r_{out}) = \alpha^3 \int_{-1}^{1} dx A_{12}(\ell_t\ell_pL_iL_{out}L_x;x) \\
\times \left\{ \Phi_{n\ell_t}(r_i(x))V_{eff}\Phi_{n\ell_p}(r_p(x)) \right\} 
\]

where

\[
x = \cos \phi = \hat{r}_i \cdot \hat{r}_{out} \tag{2.74}
\]

and \(r_t, r_p\) are functions of \(x\) through

\[
r_t(x) = [s_1^2r_i^2 + t_1^2r_{out}^2 + 2s_1t_1r_i r_{out}x]^{1/2} \\
r_p(x) = [s_2^2r_i^2 + t_2^2r_{out}^2 + 2s_2t_2r_i r_{out}x]^{1/2} \tag{2.75}
\]

The angular factor \(A_{12}\) in Eq. 2.73 is:

\[
A_{12}(\ell_t\ell_pL_iL_{out}L_x;x) = -\frac{1}{2}(-)^{1/2(L_{out}+L_i+\ell_p-\ell_t)} \\
\times [(2L_i + 1)(2L_{out} + 1)(2\ell_t + 1)(2\ell_p + 1)]^{1/2} \\
\times \sum_{M_x\mu M_x\mu} \left( \begin{array}{ccc} \ell_t & \ell_p & L_x \\ m & M_x - m & -M_x \end{array} \right) \left( \begin{array}{ccc} L_i & L_{out} & L_x \\ \mu & M_x - \mu & -M_x \end{array} \right) \Lambda_{M_x}^{\ell_p}\Lambda_{M_x}^{\ell_t}\Lambda_{M_x}^{L_i}\Lambda_{M_x}^{L_{out}} \\
\times \cos[m\phi_t + (M_x - m)\phi_p - \mu \phi] \tag{2.76}
\]

where \(\ldots\) are 3j-symbols, \(\phi\) is defined by Eq. 2.74, and

\[
\phi_t = (-)^S \cos^{-1}(\hat{r}_t \cdot \hat{r}_{out}) = (-)^S \cos^{-1}\left(\frac{s_1r_i x + t_1r_{out}}{r_t}\right) \\
\phi_p = (-)^S \cos^{-1}(\hat{r}_p \cdot \hat{r}_{out}) = (-)^S \cos^{-1}\left(\frac{s_2r_i x + t_2r_{out}}{r_p}\right) \tag{2.77}
\]

In the above, \(\phi\) and \(\cos^{-1}\) are between 0 and \(\pi\) and

\[
(-)^S = -1 \text{ (pickup)}, \quad \tag{2.78}
\]

\[
= +1 \text{ (stripping)} \quad \tag{2.79}
\]

---

*6R. Balian and E. Brézin, Nuovo Cimento 61B, 403 (1969)*
Finally, ***LINE UP *******

\[ \Lambda^q_k = 0 \]

if \( k \pm q \) is odd,

\[ = (-1)^{k+q} \frac{(k+q)!(k-q)!}{2^k(k+q)!} \]

if \( k \pm q \) is even.

### 2.3.5 DWBA Amplitude and Effective Interaction

The transfer amplitude in DWBA has the form

\[
T(\vec{k}_i \rightarrow \vec{k}_{out}) = J \int \int d^3r_i d^3r_{out} [\chi^-(k_{out}, r_{out})]^* \times \langle B, b | V_{eff} | A, a \rangle \chi^+(\vec{k}_i, \vec{r}_i) \tag{2.81}
\]

where \( \langle V_{eff} \rangle \) denotes a matrix element with respect to all internal core coordinates; \( \langle V_{eff} \rangle \) is a function of \( \vec{r}_t \) and \( \vec{r}_p \). The transition operator \( V_{eff} \) is the part of the sum of the two-body interactions between constituents of the colliding species in either channel that is not contained in the optical potential in that channel. According to Eq. 2.69 and 2.70:

\[
V_{eff} = \sum_{i \in C_t} \sum_{j \in H_t} V_{ij} - U_{opt}(r_{C_t H_t})
\]

or

\[
V_{eff} = \sum_{i \in C_p} \sum_{j \in H_p} V_{ij} - U_{opt}(r_{C_p H_p}) \tag{2.82}
\]

Two standard approximations are then made.

(i) The additional particles \( x \) have little influence on the core-core optical potential. \( U_{opt} \) thus describes the core-core interaction.

\[
U_{opt} \simeq \sum_{i \in C_t} \sum_{j \in C_i} V_{ij} \tag{2.83}
\]

and thus

\[
V_{eff} \simeq \sum_{i \in C_p} \sum_{j \in x} V_{ij} \quad \text{or} \quad V_{eff} \simeq \sum_{i \in C_t} \sum_{j \in x} V_{ij} \tag{2.84}
\]

(ii) The sum of two-body interaction \( \sum_{j \in C} \sum_{j \in x} V_{ij} \) is replaced by a one-body potential \( V(r_{C_x}) \) depending only on the relative position of the mass center of \( x \) to that of the core

\[
V_{eff} \simeq V_{C_p x}(r_p) \quad \text{or} \quad V_{eff} \simeq V_{C_t x}(r_t) \tag{2.85}
\]
2.3. TRANSFER REACTIONS

It is the second approximation that reduces x-nucleon transfer to a three-body problem. In the simplified form (Eq. 2.85) \(V_{eff}\) can be associated with one or the other vertex. For the one-body potential \(V_{C_x}\), PTOLEMY uses the potential that binds the composite system \(H\) at the appropriate vertex.

It is known that the approximation Eq. 2.83 is poor for the Coulomb part of the interaction. In this case a simple correction\(^7\) can be made;

\[
V_{eff} = V_{C_p}(r_p) + \Delta V
\]

or

\[
V_{eff} = V_{C_t}(r_t) + \Delta V
\]

\((2.86)\)

where

\[
\Delta V = V(r_{cc}) - V_{opt}
\]

\((2.87)\)

In Eq. 2.87, \(V_{opt}\) is the appropriate optical potential as indicated in the following table:

<table>
<thead>
<tr>
<th>Interaction at p vertex</th>
<th>Pick-up</th>
<th>Stripping</th>
</tr>
</thead>
<tbody>
<tr>
<td>Interaction at t vertex</td>
<td>(V_{opt}(r_i))</td>
<td>(V_{opt}(r_{out}))</td>
</tr>
</tbody>
</table>

\(V(r_{cc})\) is the optical potential between the two cores and is evaluated using the same potential parameters as \(V_{opt}\). With the Coulomb correction \(\Delta V\), \(V_{eff}\) becomes a function of \(r_p\), \(r_t\), and \(x\) in a fashion which adds no essential complication to the integral in Eq. 2.73.

2.3.6 Angular Momenta

The angular momenta transferred at the vertices, \(J_t\) and \(J_p\), have been defined in Eq. 2.59 and 2.64. The total transferred angular momentum \(L_x\) is defined by

\[
\vec{L}_x + \vec{J}_p = \vec{J}_t
\]

\((2.88)\)

The multipole or angular-momentum decomposition of the DWBA amplitude and cross section is based on the total angular-momentum transfer.

2.3.7 Differential Cross Section

\[
\frac{d\sigma}{d\Omega}(\theta) = \left(\frac{1}{E_i E_{out}}\right) \frac{k_{out}}{k_i} R \sum_{J_t J_p L_x M_x} \left|G_{M_x}^{L_x(J_t J_p)}(\theta)\right|^2
\]

\((2.89)\)

where \(R\) is a spin-statistical factor:

\[
R = \frac{2J_b + 1}{2J_a + 1} \text{ (pick – up)}
\]

\[
= \frac{2J_B + 1}{2J_A + 1} \text{ (stripping)}
\]

\((2.90)\)

In PTOLEMY only one value of \( J_t \) and \( J_p \) is allowed and the sum over \( J_t \) and \( J_p \) in Eq. 2.89 consists of only one term.

The multipole transition amplitude \( G \) is a sum of products of spectroscopic and geometrical components

\[
G^{L_x(J_tJ_p)}_{M_x} (\theta) = \sum_{n_t\ell_t n_p\ell_p} A_{L_xJ_tJ_p}(n_t\ell_t n_p\ell_p) B^{L_x(J_tJ_p)}_{M_x}(n_t\ell_t n_p\ell_p; \theta) \tag{2.91}
\]

summed over all contributing states of orbital motion of the transferred nucleons at each vertex. PTOLEMY allows only one projectile and target bound state so the sum in Eq. 2.91 consists of only one term. With the phase convention introduced earlier, \( G \) is the negative of the amplitude used in Schwarzchild, et al.

### 2.3.8 Geometrical Component (including radial integrals)

The angular dependence of the cross section is contained in the “geometrical” component of Eq. 2.91:

\[
B^{L_x(J_tJ_p)}_{M_x}(n_t\ell_t n_p\ell_p; \theta) = \sqrt{4\pi \ell_x + \ell_t - \ell_p} \sum_{L_i L_{out}} C^{L_{out}L_xL_i}_{M_x,-M_x,0} I_{L_i L_{out}L_x} (n_t\ell_t n_p\ell_p) \sqrt{L_{out}} (\theta) \tag{2.92}
\]

The radial integrals are expressed in terms of the bound-state form factor \( H \) [Eq. 2.73 to 2.80] and the radial scattering functions by

\[
I_{L_i L_{out}L_x} (n_t\ell_t n_p\ell_p) = e^{i(\sigma_i + \sigma_{out})} \int \int r_i dr_i r_{out} dr_{out} f_{L_i} (r_i) H_{L_i L_{out}L_x} (n_t\ell_t n_p\ell_p) f_{L_{out}} (r_{out}) \tag{2.93}
\]

Note that \( B^{L_x}_{M_x} \) is independent of the shell-model wave functions of the nuclear states.

### 2.3.9 Spectroscopic Factors and Spectroscopic Component

Let the states \( A, a, B, b \) be represented by shell-model wave functions. In order to treat the center-of-mass variable \( \vec{R} \) consistently and to separate internal and center-of-mass variables of the nuclear states, we use harmonic-oscillator shell-model wave functions with the center-of-mass motion in its ground (0s) state.

Consider then the internal states at the target vertex with \( J_t \) the total angular momentum transfer at that vertex. Let \( \psi^\gamma_{J_t} \) be a complete set of \( x \)-nucleon shell-model states for the transferred nucleons, and let \( a^+ (\gamma_t J_t M_t) \) be the creation operators that produce these states from the vacuum. The \( x \)-nucleon states must be projected onto states of the transferred nucleons of the form Eq. 2.57, with internal and center-of-mass variables separated. Define the necessary coefficients

\[
K_{J_t}(n_t\ell_t x J_x; \gamma_t) = \\
\int \int d^3 R_x d^3 \rho_x ([\psi^{n_t\ell_t}(\vec{R}_x) \times \phi^{J_x}(\rho_x)]_{M_t})^* \psi_{M_t}^\gamma (\{\vec{r}_x\}) \tag{2.94}
\]
where \( \{ \vec{r}_\alpha \} \) is a set of nucleon coordinates and

\[
R_x = \frac{1}{x} \sum_{\alpha=1}^{x} \vec{r}_\alpha
\]

Then the conventional spectroscopic amplitude \( \sqrt{S} \) is given by

\[
\sqrt{S(n_t \ell_t, xJ_x, J_T; H_t, C_t)} = \sum_{\gamma_t} K_{J_t}(n_t \ell_t, xJ_x; \gamma_t) \langle J(H_t) | A^{\dagger} | J(C_t) \rangle
\]

(The reduced matrix element is defined by the Wigner-Eckart theorem in the form

\[
\langle jm | T^k_{q} | j'm' \rangle = C_{m'qm}^{j'kj} \langle j | T^k | j' \rangle.
\]

The treatment of the centers of mass leading to Eq. 2.94 and 2.96 is exact if the shell-model wave functions and the radial functions \( \psi \) in 2.56 and its projectile analog are harmonic-oscillator functions. However, the radial functions of the target and projectile bound states are eigenfunctions of Woods-Saxon potentials. This difficulty is usually ignored since the level of precision of the entire analysis (in particular its absolute normalization) is seldom high enough to require consideration of such niceties. A crude correction factor can be introduced by expanding the Woods-Saxon wave functions in terms of oscillator functions and assuming that one term dominates. It can then be shown that the replacement

\[
\sqrt{S(n_t \ell_t, xJ_x, J_T; H_t, C_t)} \rightarrow \Theta(n_t \ell_t, xJ_x, J_T; H_t, C_t)
\]

\[
\Theta(H_t, C_t) = \left( \frac{H_t}{C_t} \right)^{2n_t + \ell_t} \frac{2L_x + 1}{2} \sqrt{S(H_t, C_t)}
\]

should be correct for the use of the oscillator shell-model wave functions. The projectile vertex is handled in the same way.

In Ptolemy, the spectroscopic amplitudes \( \Theta \) are read in directly; they can often be inferred from suitable light-ion reactions between the nuclear state in question. Note the \( S \) as defined above reduces to the standard spectroscopic factor in the case of single-nucleon transfer.

The spectroscopic component \( A \) [Eq. 2.91] is given in terms of the spectroscopic amplitudes \( \Theta \) [Eq. 2.96 and 2.97] by

\[
A_{L_xJ_tJ_p}(n_t \ell_t n_p \ell_p) = \sqrt{2L_x + 1} \sum_{xJ_x} \left(-1\right)^{J_x - J_p + \ell_p + \ell_t} W(\ell_t, J_t \ell_p, J_p; J_x L_x) \Theta(n_t \ell_t, xJ_x, J_t; H_t C_t) \Theta(n_p \ell_p, xJ_x, J_p; H_p C_p)
\]

\( A \) is independent of the scattering angle and \( M_x \).

### 2.3.10 Outline of Steps in a DWBA Computation for Transfer Reactions

The main steps in a DWBA transfer reaction calculation can be schematically summarized as follows. In practice a number of these steps are carried out in parallel.
1. Adjust the potentials at the interaction vertices to reproduce the experimental separation energies and compute the bound-state wave functions. This specifies the effective transition operator through Eq. 2.85 or 2.86.

2. For given optical-model parameters, solve Eq. 2.27 – 2.29 for the radial scattering functions. At this stage elastic-scattering amplitudes and cross sections can also be computed.

3. Use Eq. 2.73 to 2.80 to compute the bound-state form factors $H(r_i, r_{out})$.

4. Fold the bound-state form factors with the radial scattering functions and integrate Eq. 2.93 to obtain the radial integrals $I_{L_iL_{out}L_x}$.

5. Using given spectroscopic amplitudes $\Theta$ compute the spectroscopic components $A_{L_x}$ of the multiple amplitudes using Eq. 2.98.

6. Calculate the geometrical components $B_{M_x}^{L_x}$ of the multipole amplitudes using Eq. 2.92.

7. Construct the multipole components of the transition amplitude using Eq. 2.91 and compute the cross section [Eq. 2.89].

Note that in heavy-ion calculations, more than 90% of the time is spent carrying out steps (3) and (4) – construction of the bound-state form factors and integration over $r_i$ and $r_{out}$.

### 2.4 Constants and Units

PTOLEMY uses the values:\(^8\)

\[
\hbar c = 197.32858 \text{ MeV} \cdot \text{fm} \tag{2.99}
\]

\[
M_\mu = 931.5016 \text{ MeV}/c^2 \tag{2.100}
\]

\[
\alpha^{-1} = 137.03604 \tag{2.101}
\]

where $M_\mu$ is the atomic mass unit and $\alpha$ is the fine-structure constant.

PTOLEMY reads, stores, and prints quantities in the following units:

### 2.4. CONSTANTS AND UNITS

<table>
<thead>
<tr>
<th>Quantity</th>
<th>Unit</th>
</tr>
</thead>
<tbody>
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<td>degrees (input, output)</td>
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<tr>
<td></td>
<td>degrees or radians (internal)</td>
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<tr>
<td>cross sections</td>
<td>mb</td>
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<tr>
<td>lengths, radii</td>
<td>fm</td>
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<td>MeV/c$^2$</td>
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<tr>
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<tr>
<td>potentials</td>
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<tr>
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</tr>
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<td>elastic amplitudes ($F, B$)</td>
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</tr>
<tr>
<td>amplitudes ($G, B$)</td>
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<tr>
<td>radial integrals ($I$)</td>
<td>MeV $\cdot$ fm</td>
</tr>
<tr>
<td>transfer form factors ($H$)</td>
<td>MeV $\cdot$ fm$^{-3}$</td>
</tr>
<tr>
<td>inelastic effective interaction ($H$)</td>
<td>MeV</td>
</tr>
</tbody>
</table>

Note that since cross sections are expressed in mb, a factor of 10 is necessary in Eq. 2.12 – 2.14, 2.23 – 2.26, 2.35, and 2.89 to convert from fm$^2$. 
Chapter 3

Notation and Syntax

3.1 Notation

Ptolemy ignores the case of input letters; lowercase letters are converted to uppercase before further processing. Thus some changes must be made to the notation established in the previous section for the description of inelastic excitation and transfer reactions. In general lower case letters will simply be converted to upper case. However, in Chapter 2 a distinction was made between lower- and upper-case letters in the identification of the reaction participants. For PTOLEMY input this distinction will be maintained by specifying the target particles as A and B while the projectile particles will be a and b. Thus the reaction computed by PTOLEMY may be written as

\[ A(a,b)B \]

In the incoming state, the target is referred to as \( A \) and the projectile is \( A \). In the final state the residual target is \( B \) and the ejectile is \( b \). The exchanged particle is referred to as \( X \). For a stripping reaction we have

\[
\begin{align*}
    a &= b + x \\
    B &= A + X
\end{align*}
\]

while for a pickup reaction

\[
\begin{align*}
    b &= a + x \\
    A &= A + X
\end{align*}
\]

PTOLEMY will compute either pickup or stripping reactions; it is not necessary for the user to interchange particles to force the reaction into one form or the other. Inelastic excitation is implied if the mass and charge of \( a \) and \( b \), and of \( A \) and \( B \), are the same.

The projectile or ejectile bound state (whichever is appropriate) is always referred to as the projectile bound state while the target or residual nucleus bound state is called the target bound state. The \( Q \) value of the reaction is the difference of the outgoing and incoming kinetic energies in the center of mass system so that in terms of the bound state energies

\[
\begin{align*}
    Q &= E(\text{projectile}) - E(\text{target}) : \text{stripping} \\
    Q &= E(\text{target}) - E(\text{projectile}) : \text{pickup}
\end{align*}
\]

Note that PTOLEMY deals with actual bound state energies (i.e., negative numbers). In the case of inelastic excitation, the \( Q \)-value is, of course, just the negative of the
excitation energy.

3.2 Ptolemy Syntax

PTOLEMY uses a free-form keyword-based input. Options are specified and stages of the calculation selected by the specification of the appropriate keyword. Numeric values are entered by the specification of a number entering keyword followed by the desired value. One or more keywords and associated numbers may be included on a single input line or a data value may be on the line following its keyword. (The CHANNEL, REACTION, and HEADER keywords are exceptions and require associated information to be on the same input line.) Words and numbers may not be split across two lines and they may not contain embedded blanks. Input lines may be up to **** characters long.

Keywords may be separated from other keywords on the same line by blanks, commas, or sequences of blanks and commas. The equal sign may be used (but is not required) between a keyword and its associated data value. The colon may be used following the CHANNEL, REACTION, and HEADER keywords but should not otherwise be used in PTOLEMY input. The semicolon is used to begin a stage of the calculation; it indicates that all input needed for that stage has been provided.

Numerical data may be entered with or without a decimal point and may have the E form of exponent. Valid numerical inputs are

2,2.3,0.0002,2E-4, 2.325E+7, -5.3, +7E-5

An E appearing in a number indicates the beginning of the power of 10 by which the number is to be multiplied. Thus

5.3E-7 = 5.3 \times 10^{-7}

1E20 = 10^{+20}

Angular momenta that have the possibility of being half-integer (J or S values but not L values) have a special form of input. They may be either simple integers or integers followed by /2 to indicate half integer values. They should not be coded with a decimal point. Thus

J=2, S=3/2, JP=4/2

are all valid (the last is the same as JP=2) while

J=2., S=1.5

are both invalid. Such J and S values may be followed by a parity sign. Thus

J=2, J=2-, and J=2+

are all valid.

Keywords may have more than eight characters in their names but only the first eight characters are used and required. Keywords never have embedded blanks in their names.

Comments may be placed anywhere in the input. They are preceded by a dollar sign ($) which indicates that the rest of the input line is a comment. If a second dollar sign appears on the same line, the comment is terminated and the remainder of the line is processed as normal input.
Chapter 4

Elastic and Bound State Calculations

The basic ingredients of the DWBA calculations performed by PTOLEMY are two-body wavefunctions – optical-model scattering states for inelastic excitation and both bound-state and optical-model scattering states for transfer. Obviously, optical-model scattering states are also used in the optical model fitter.

One may use the facilities of PTOLEMY to compute the properties of two-body bound or scattering states without doing a larger scale calculation. Such calculations will be referred to as “stand–alone” two-body calculations. Wherever possible the same keywords and conventions are used for defining the two-body states that are components of a larger calculation as are used in stand-alone two-body calculations. Examples are the definitions of the potentials and the specification of the integration grid used to solve the Schroedinger equation. For this reason we describe the stand–alone two–body calculations before progressing to the more complicated calculations — the material introduced in this chapter will be referred to in many of the succeeding chapters.

4.1 Specifying the Two-Body Channel

Stand-alone two-body calculations are done by defining the two particles and the potential that acts between them and then entering a semicolon to start the calculation. If the energy is negative a bound state calculation will be made, if it is positive the two-body scattering will be computed. If desired one may use the keywords BOUNDSTATE or SCATTERING to indicate which is to be done and the energy will be checked for validity. Although it is possible to mix stand-alone and DWBA calculations in one job, it is recommended that separate jobs be used for stand-alone and complete DWBA runs. However many stand-alone calculations of both bound states and scattering may be done in one job.

The two particles involved in the stand-alone calculation are referred to as the “projectile” and “target.” These words have their customary meanings for scattering states; the only distinction for bound states is that the projectile’s angular momentum is used to determine the spin-orbit force.
4.1. SPECIFYING THE TWO-BODY CHANNEL

The CHANNEL keyword may be used to specify the nuclei in the two-body state. Some examples will illustrate its form:

CHANNEL 12C + 208PB

CHANNEL: P + 11B = C12

CHANNEL = 209BI(7/2 -- 0.9) = P + 208PB

The first example gives a scattering state while the other two define bound states. Note that in the case of bound state channels, the resultant bound state may be either the first nucleus (in which case it is followed by an equal sign) or the last nucleus. In all cases the projectile and target must be separated by a plus sign and the projectile always comes first. If a bound state is being specified, the composite nucleus may be either first or last and must be separated from the other two nuclei by an equal sign.

The nuclides are defined by an element symbol consisting of the atomic weight and a one- or two-character element abbreviation. The atomic weight may either precede or follow the symbol but no blank spaces or other punctuation may intervene. In addition the following symbols (without atomic weights) may be used:

N - neutron
P - proton
D - deuteron
T - triton
H - ^3He
A - ^4He

Excited states may be indicated by enclosing the spin and excitation energy in parentheses following the element symbol. The left parenthesis for excited state specification must immediately follow the element symbol. The excitation energy and spin of the excited state may be given in either order, and the excitation energy must include a decimal point even if it happens to be an integer. Excitation energies are given in MeV. Any or all of the two or three nuclei may be given an excited state specification. The CHANNEL keyword and the complete channel specification must be contained on a single input line.

The CHANNEL keyword will define the projectile, target and bound-state mass, charge, and intrinsic spin. The intrinsic spins and the ground-state mass excesses of the nuclei are found from the ***FIX***1975 Oak Ridge Atomic Mass Adjustment\(^1\) and the 1971 Nuclear Wallet Cards compilation. In addition for bound states the total angular momentum and the bound-state (cluster separation) energy are also defined.

If the CHANNEL keyword is not used, the particle masses may be entered with the keywords MP and MT which give the masses in AMU of the projectile and target respectively.

---
\(^1\)Brookhaven
The masses do not need to be integers. Alternatively one can use the keyword \( M \) to enter the reduced mass in MeV/c\(^2\). The charges of the two particles may be entered with the keywords \( ZP \) and \( ZT \). The excitation energies may be entered using the keywords \( E^*P \) and \( E^*T \); if they are not entered, zero will be used. The projectile and target intrinsic spins may be entered with the keywords \( SP \) and \( ST \). The total angular momentum of a bound state may be entered with the keyword \( J \). It is not necessary to enter \( J \) or \( ST \) (the calculation does not depend on them) and \( SP \) is necessary only if there is a spin-orbit force.

The projectile and target will be recognized as identical if they have the same mass, charge, spin, and excitation energy. In such cases the appropriate spin statistics will be used for scattering calculations. Non-identical particle scattering may be forced by specifying a small excitation energy for one of the particles.

The c.m. energy may be entered by using either of the keywords \( ECM \) or \( E \). In the case of bound states one of these should be used to enter the energy as a negative number unless the \textsc{channel} keyword is used. The laboratory scattering energy may be entered with the keyword \textsc{elab}. In this case both \textsc{mp} and \textsc{mt} must be defined to allow the conversion to the c.m. energy.

The number of nodes and orbital angular momenta of bound states must be defined for bound state calculations. The keyword \textsc{nodes} is used to specify the number of nodes. The node at the origin is not included in the count so that the lowest bound state for each value of \( L \) has 0 nodes. The keyword \( L \) is used to specify the orbital angular momentum of the bound state. If a spin-orbit force is being used in the bound state, it is necessary to enter the total angular momentum of the “projectile”. This is done with the keyword \textsc{jp}. \( J_p \), \( L \) and \( S_p \) are used to find the value of \( L \cdot S \) in the spin-orbit force. If either of \( S_t \) (the target spin) or \( J \) (the total bound state spin) are zero, then \( J_p \) need not be specified since it will be uniquely determined by other known spins.

### 4.2 Specifying the Potentials

The potentials are defined by the keywords \( V_i \), \( R_i \) or \( R_i0 \), and \( A_i \) where the suffix “\( i \)” indicates the potential that is being defined. Possibilities for “\( i \)” are:

- \texttt{null} – no suffix refers to the real part of the Woods-Saxon well.
- \texttt{I} – The suffix \texttt{I} refers to the imaginary part of the Woods-Saxon well.
- \texttt{S0} – The suffix \texttt{S0} designates the real part of the spin-orbit force.
- \texttt{S0I} – Imaginary part of the spin-orbit force.
- \texttt{SI} – Imaginary surface potential.
- \texttt{C} – Coulomb potential (\texttt{VC} and \texttt{AC} are not defined).

The forms of these potentials are as follows:
4.2. SPECIFYING THE POTENTIALS

1. Real part of the Woods-Saxon:

\[
\frac{-V}{(1 + X)},
\]

\[X = \exp \left[ \frac{(r - R)}{A} \right].\] (4.1)

2. Imaginary part of the Woods-Saxon (volume absorption):

\[
\frac{-VI}{1 + X_I},
\]

\[X_I = \exp \left[ \frac{(r - RI)}{AI} \right].\] (4.3)

3. Real part of the spin-orbit:

\[+ (VSO + \text{T} \times V) \times 4L \cdot S \times \frac{1}{r} \times \frac{d}{dr} \frac{1}{(1 + X_{SO})},\] (4.5)

\[X_{SO} = \exp \left[ \frac{(r - RSO)}{ASO} \right].\] (4.6)

4. Imaginary part of the spin-orbit:

\[+ (VSOI + \text{T} \times VI) \times 4L \cdot S \times \frac{1}{r} \times \frac{d}{dr} \frac{1}{(1 + X_{SOI})},\] (4.7)

\[X_{SOI} = \exp \left[ \frac{(r - RSI)}{ASI} \right].\] (4.8)

5. Imaginary surface potential (surface absorption):

\[+ VSI \times 4ASI \times \frac{d}{dr} \frac{1}{(1 + X_{SI})},\] (4.9)

\[X_{SI} = \exp \left[ \frac{(r - RSI)}{ASI} \right].\] (4.10)

6. Coulomb potential

(a) point and uniform sphere:

\[
\frac{+Z_pZ_ie^2}{r} : r \geq RC,
\]

\[
+Z_pZ_ie^2 \times \left\{ 3 - \left( \frac{r}{RC} \right)^2 \right\} \quad : \quad r < RC,
\] (4.11)

\[\frac{(2RC)}{\left( \frac{r}{2RC} \right)^2} \]
(b) two uniform spheres:

$$\frac{9Z_pZ_te^2}{(16\pi^2 RCP^2 RCT^2)} \times \int_0^{RCP} d^3r_p \int_0^{RCT} d^3r_t \times \frac{1}{|r - r_p - r_t|},$$

(4.12)

where

$$L \cdot S = \frac{(J_p(J_p + 1) - L(L + 1) - S_p(S_p + 1))}{2}.$$

(4.13)

The potential well depths are given in MeV. Note that the spin-orbit well depths may either be specified directly by using the $VSO$ and $VSOI$ keywords or their ratio to the corresponding Woods-Saxon depths may be given by using the $TAU$ and $TAUI$ keywords. The $TAUs$ are related to the $LAMBDA$s of DWUCK and LOLA by

$$TAU = \frac{\text{LAMBDA}}{(4 \times 45.2)}.$$

(4.14)

Note that $VSO$ and $VSOI$ have dimensions of MeV since the factor “4” in the definition of the spin-orbit force is interpreted as $2 \times 2$ where one “2” converts $L \cdot S$ to $L \cdot \sigma$, and the other “2” is approximately the square of the pion Compton wavelength in fm. The ratio $TAU$ is dimensionless. The spin-orbit force always refers to the spin of the projectile coupled to its orbital angular momentum; the spin of the target does not enter the potential. A spin-orbit force may not be used if the particles are identical.

In all cases the radius parameter ($R_0$, $R_{IO}$, $RSO_0$, $RSOI_0$, $RSI_0$, $RC_0$, $RC0P$, or $RC0T$) may be entered in place of the potential radius. The radius is then computed as

$$R = R_0 \times \frac{1}{M_1^{1/3}} : M_p \leq 2 \text{ or } RO\text{TARGET specified},$$

$$R = R_0 \times \left\{ \frac{1}{M_1^{1/3}} + \frac{1}{M_t^{5/3}} \right\} : M_p \geq 3.$$

(4.15)

These formulas are generally appropriate for optical potentials, but may result in unexpectedly large potential radii for bound states of a nucleon cluster (such as an alpha particle) and a heavier core. As noted, the keyword $RO\text{TARGET}$ may be used to cause all subsequent radius parameters to be computed using only $M_t^{5/3}$.

Defining $RC$ or $RC0$ causes the “point and uniform sphere” Coulomb potential (6a) to be used; otherwise the “two uniform spheres” (6b) potential is used. In the latter case, the radius of each nucleus may be explicitly entered with the $RCP$ and $RCT$ or $RC0P$ and $RC0T$ keywords, or they may be left undefined. If they are not defined, $\text{PTOLEMY}$ will choose them to give RMS radii roughly consistent with those determined by electron scattering. These radii are taken to be:

$$1 \leq A \leq 2 \quad R = 0,$$

$$3 \leq A \leq 16 \quad \text{Values from Landolt-Boernstein}^2,$$

$$17 \leq A \quad \text{Values from Bohr and Mottelson formula}^3.$$

The Coulomb potential between two uniform spheres is not available for bound states.

The real and imaginary potential parameters may be given a dependence on the laboratory energy by the use of keywords that end in $E$ or $ESQ$. In this case the quantities to be used in the above equations will be computed as follows:
4.3. CONTROLLING THE CALCULATION

\[ a = A + AE \times E_{\text{lab}} + AESQ \times E_{\text{lab}}^2 \]
\[ R_O = RO + ROE \times E_{\text{lab}} + ROESQ \times E_{\text{lab}}^2 \]
\[ V = V + VE \times E_{\text{lab}} + VESQ \times E_{\text{lab}}^2 \]
\[ a_i = AI + AIE \times E_{\text{lab}} + AIESQ \times E_{\text{lab}}^2 \]
\[ R_iO = RIO + RIOE \times E_{\text{lab}} + RIOESQ \times E_{\text{lab}}^2 \]
\[ V_i = VI + VIE \times E_{\text{lab}} + VIESQ \times E_{\text{lab}}^2 \]

The names appearing on the right of the equal signs in the above equations are the keyword values the user enters. The quantities on the left are then used to evaluate the potentials. The default value for all the keywords ending in \( E \) or \( ESQ \) is 0.

The keyword \texttt{EINVERSE} may be used to indicate that \( \frac{1}{E_{\text{lab}}} \) and \( \frac{1}{E_{\text{lab}}^2} \) are to be used in the above formulas for the energy dependent parameters. The default is \texttt{EPowers} which results in the above formulas.

If \( Ai \) and/or both \( Ri \) and \( Rf0 \) are not defined for the imaginary part of a potential, \texttt{PTOLEMY} will use the \( Ai \) or \( Ri \) for the real part of the same potential (\( RS0 \) will be used for \( RS0I \), \( A \) for \( AI \), etc.). If \( AS0 \) or both \( RS0 \) and \( RS00 \) are not defined, \( A \) or \( R \) will be used for them. If \( ASI \) or both \( RSI \) and \( RSI0 \) are not defined, \( AI \) or \( RII \) will be used for them.

In process of computing the bound state wavefunctions it is necessary that \( V \) (the potential depth) and \( E \) (the bound state energy) be made consistent with each other. \texttt{PTOLEMY} varies one or the other of these two quantities until they are consistent. The keywords \texttt{FITV} and \texttt{FITE} may be used to determine which quantity is to be varied. \texttt{FITV} causes \( V \) to be changed to produce a well that has the bound state energy \( E \). If a spin-orbit force has been specified via the keyword \texttt{TAU}, the depth of the spin-orbit force is also varied since the ratio of the spin-orbit force to the Woods-Saxon well is held constant at \( TAU \). On the other hand, if \( VSO \) is used to specify the spin-orbit force, the strength of the spin-orbit force is not changed as \( V \) is changed. \texttt{FITE} causes \( E \) to be computed as the bound state energy of the given potential. The default is \texttt{FITV}.

4.3 Controlling the Calculation

If a scattering calculation is being made, the S-matrix elements will be found for a range of orbital angular momentum values. This range may be explicitly specified by the \texttt{LMIN} and \texttt{LMAX} keywords. If \texttt{LMIN} and/or \texttt{LMAX} are not specified, they will be based on \( L_c \) (the angular momentum for which \(|S| = 1/2\)) which is estimated via semi-classical expressions. In such cases the four keywords \texttt{LMINMULT}, \texttt{LMINSUB}, \texttt{LMAXMULT}, and \texttt{LMAXADD} are used to compute \texttt{LMIN} and/or \texttt{LMAX} from the formulas:

\[
\text{LMIN} = \min(\text{LMINMULT} \times L_c, L_c - \text{LMINSUB})
\]
\[
\text{LMAX} = \max(\text{LMAXMULT} \times L_c, L_c - \text{LMAXADD})
\]

Note that if only one of \texttt{LMIN} or \texttt{LMAX} is explicitly specified, then only the other is computed from the above formulas. One should note that the extrapolation to large \( L \)-values that is provided in DWBA calculations does not occur in elastic scattering calculations. Thus a larger \texttt{LMAX} or \texttt{LMAXADD} is required for elastic scattering calculations.

It may be desired to compute the elastic \( S \)-matrix elements for only one value of \( L \). In such cases, the keyword \texttt{L} should be used to specify the desired value. If a spin-orbit force is entered for a scattering problem, the \( S \)-matrices will be computed for all values
of $Jp$ connected with each value of $L$ (note that $Sp$ is not limited to $1/2$ for spin-orbit forces). If it is desired to have only one value of $Jp$, the keyword $JP$ may be used to specify that value. If both of the keywords $L$ and $JP$ are used, only one scattering partial wave will be computed. Bound states are always computed for only one value of $L$ and $Jp$.

The keyword $\text{ELASTIC}$ may be used to cause the elastic differential cross sections to be computed. The default is $\text{NOELASTIC}$ which suppresses the differential cross sections. The cross sections are given in millibarns/steradian and as ratios to the corresponding Rutherford (Mott if the particles are identical) cross sections. The cross sections are averaged over the initial spins and summed over the final spins.

The grid of c.m. angles on which the differential cross sections are displayed is controlled by the keywords $\text{ANGLEMIN}$, $\text{ANGLEMAX}$, and $\text{ANGLESTEP}$. The angles are given in degrees. The default values are

\begin{align*}
\text{ANGLEMIN} &= 0, \\
\text{ANGLEMAX} &= 90, \\
\text{ANGLESTEP} &= 1.
\end{align*}

The keyword $\text{LABANGLES}$ may be used to indicate that $\text{ANGLEMIN}$, $\text{ANGLEMAX}$, and $\text{ANGLESTEP}$ specify a grid of laboratory angles. In such cases there is a two to one mapping of laboratory angles to c.m. angles if the projectile mass is greater than the target mass. $\text{PTOLEMY}$ will convert positive laboratory angles to the smaller c.m. angle and negative angles to the larger c.m. angle. In such cases $\text{ANGLEMAX}$ may be negative to cause the c.m. angles to steadily increase through $90^\circ$. The default is $\text{CMANGLES}$.

The computation of the two-body wavefunctions (both bound and scattering states) may be controlled with the $\text{ASYMPTOPIA}$ and $\text{STEPSIZE}$ or $\text{STEPSPER}$ keywords. $\text{ASYMPTOPIA}$ specifies (in fm) the radius at which the wavefunctions are to be assumed to be asymptotic. It is also the largest value of $r$ for which the wavefunctions will be computed and stored.

The keyword $\text{STEPSIZE}$ gives the increment used in the solution of the bound- and scattering-state Schrödinger equations. Since arrays must be constructed that have $\text{ASYMPTOPIA}/\text{STEPSIZE}$ elements, one should avoid making this ratio very large. The keyword $\text{STEPSPER}$ may be used to specify the number of steps to use per wavelength. If it is entered, $\text{STEPSIZE}$ will be computed according to the formulas:

\begin{align*}
\text{STEPSIZE} &= \min \left( \frac{\kappa}{a}, \frac{\lambda}{2 \times a} \right) / \text{STEPSPER} : \text{Bound states} \\
\text{STEPSIZE} &= \min(\lambda, 2 \times a) / \text{STEPSPER} : \text{Scattering}
\end{align*}

where $\kappa$ is the bound state inverse range:

\begin{equation}
\kappa = \sqrt{2M |\zeta|} \quad (4.16)
\end{equation}

and $\lambda$ is the scattering wavelength:

\begin{equation}
\lambda = \frac{2\pi}{\sqrt{2ME}}. \quad (4.17)
\end{equation}

In both cases $a$ is the diffuseness of the real part of the central Woods-Saxon well. It is suggested that $\text{STEPSPER}$ be used instead of $\text{STEPSIZE}$ since then the step size will
4.4. THE TWO-BODY WAVEFUNCTIONS

<table>
<thead>
<tr>
<th>Keyword</th>
<th>Default</th>
<th>EL1</th>
<th>EL2</th>
<th>EL3</th>
</tr>
</thead>
<tbody>
<tr>
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<td>25</td>
</tr>
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<td>$2\times10^{-4}$</td>
<td>$2\times10^{-5}$</td>
</tr>
</tbody>
</table>

Table 4.1: PARAMETERSET names and associated values for elastic scattering calculations. The first column gives the default values.

automatically be adjusted as the wavelength changes due to changes in the scattering energy. If both STEPSIZE and STEPSPER are defined, STEPSPER has precedence.

The keyword PARAMETERSET may be used to select a standard set of values for the calculation-controlling keywords. The keyword is followed by the name of the desired set; Table 4.3 gives the names of the sets for elastic scattering and the associated values.

Individual settings may then be overridden by subsequently entering the appropriate keywords. Note that STEPSPER and not STEPSIZE is defined by these PARAMETERSET sets. Since STEPSPER has precedence over STEPSIZE, one must use the command

UNDEFINE STEPSPER, STEPSIZE=ssss

if one wants to enter a specific STEPSIZE after having used PARAMETERSET (UNDEFINE is defined in Chapter 8). The FITACCURACY keyword in Table 4.3 will be explained in the chapter on optical model fits.

4.4 The Two-Body Wavefunctions

The computed wave functions (for both bound and scattering states) will be printed if the keyword WRITESTEP is used. This keyword specifies the stepsize for which the wavefunction is to be tabulated. The value of WRITESTEP should be a multiple of STEPSIZE; if it is not, the closest multiple of STEPSIZE will be used. Setting WRITESTEP equal to 0 (the default) will suppress the printing of the wavefunction.

The bound state wavefunctions are the solutions of the Schrödinger equation

$$\left\{ \frac{\hbar^2}{(2Mr^2)} \left[ - \frac{d}{dr} \right] r^2 \left( \frac{d}{dr} \right) + L(L+1) \right\} \Phi(r) = 0, \quad (4.18)$$

while the scattering wavefunctions are the solutions of

$$\left\{ \frac{\hbar^2}{(2M)} \left[ - \frac{d^2}{dr^2} + \frac{L(L+1)}{r^2} \right] + V(r) - E \right\} f_L(r) = 0. \quad (4.19)$$

The bound state wavefunctions are normalized to unity so that

$$\int_0^\infty dr r^2 \Phi^2 = 1 \quad (4.20)$$
The scattering wavefunctions are normalized to have the asymptotic form

\[ f(r) \rightarrow \frac{1}{2} \left\{ (1 + S)F(kr) + i(1 - S)G(kr) \right\}, \]

\[ = \cos(\delta) \exp(i\delta) \left\{ F(kr) + \tan(\delta)G(kr) \right\}, \tag{4.21} \]

where \( F \) and \( G \) are the regular and irregular Coulomb functions, and \( \delta \) is the complex phase shift \( S = \exp(2i\delta) \). If the optical potential is real, then the phase shift \( \delta \) is real and the phase of the wavefunction is \( \exp(i\delta) \) for all \( r \). In this case one might want to use the REALWAVE keyword to cause the wavefunctions to be multiplied by \( \exp(-i\delta) \) so that

\[ f(r) \rightarrow \cos(\delta)F(kr) + \sin(\delta)G(kr) \quad \text{REALWAVE}. \tag{4.22} \]

The default is COMPLEXWAVE. Note that asymptotically the bound state wave functions behave as \( \exp(-r) \) while the scattering wavefunctions do not have a \( 1/r \) in their asymptotic form.

The name of the bound state wavefunction will be \( \text{PHIn} \) where “n” is an integer that is 1 for the first bound state and is increased by 1 for each subsequent bound state. After \( n = 9 \), it is set back to 1 again. Thus one can have up to nine bound state wavefunctions in the allocator at once. The names of the real and imaginary scattering wave functions will be \( \text{WAVER} \) and \( \text{WAVEI} \). If it is desired to have more than one scattering wavefunction in the allocator at once, the keywords RENAME or COPY should be used (see Chapter 8).

The keyword CHECKASYMPT may be used to cause the rate of convergence of the scattering wavefunctions to the asymptotic form given above to be displayed when the wavefunction is computed. The difference of the exact wavefunction and the asymptotic form will be printed at intervals determined by WRITESTEP (which must also be defined). The keyword NOCHECKASYMPT cancels a previously entered CHECKASYMPT and is the default.

### 4.5 Reading the Output

PTOLEMY produces a summary of the two-body channel that for the most part is self-explanatory. The summary contains properties of the nuclei (spin, mass, etc.) and the two-body scattering or bound-state energy. Following this is the potential summary in which there is a line for each non-zero potential. The well depths are given in a column labeled “Coupling Cons.”; the entry in this column for the Coulomb potential is the Sommerfeld parameter. If the calculation is of a bound state, the summary is produced after the calculation is complete and thus contains the real well depth (or binding energy if \text{FITE} \ was specified) that is the result of the search for an eigenvalue of the Schrödinger equation.

During the computation of elastic wavefunctions, the S-matrix elements, their magnitudes and phases (in degrees), and the transmission coefficients are tabulated. If the keyword ELASTIC is specified, the tabulation of S-matrix elements is followed by a tabulation of elastic cross sections. In this tabulation scattering angles (in degrees) and differential cross sections are given in both the c.m. and laboratory frames. In addition the c.m. Rutherford cross section and the ratio of the elastic cross section to the Rutherford value are given. If the scattering is of identical particles, then all of the cross...
sections are suitably symmetrized (the resulting symmetrized Rutherford cross sections are sometimes referred to as Mott cross sections).

The last two columns of this tabulation are labeled “% PER LOW L” and “% PER HIGH L” and contain indications of the errors in the differential cross sections due to the low and high orbital angular momentum limits. They are defined as

\[
% \text{PER LOW } L = 100 \left( \frac{\sigma(L_{\text{MIN}}) - \sigma(L_{\text{MIN}} + 2)}{2} \right)
\]

\[
% \text{PER HIGH } L = 100 \left( \frac{\sigma(L_{\text{MAX}}) - \sigma(L_{\text{MAX}} - 2)}{2} \right)
\]

and thus are the percent error for omitting the smallest and largest \(L\)-value used in the calculation. Experience has shown that the actual error in the cross sections is typically five times the number printed, but such an estimate is strongly dependent on the rate at which \(|S|\) is approaching 0 or 1. (Of course, if \(L_{\text{MIN}} = 0\) the low \(L\) error may be ignored.)

Following the differential cross section tabulation, the total reaction cross section and the nuclear total cross section are printed. These quantities are defined in Chapter 2.
Chapter 5

Optical Model Potential Fits

PTOLEMY provides a powerful and efficient program for fitting optical model potentials to elastic scattering data. Data for more than one elastic scattering reaction or at more than one bombarding energy may be used in a fit. The optical potentials may be given an energy dependence. The normalization of the data and the laboratory angle calibration may be used as fit parameters.

5.1 Specifying the Fit Parameters and Data

The input for an optical model fit consists of the following items:

1. specification of the potential parameters that are to be varied in the fit,

2. initial values of the search parameters and the fixed values of all other potential parameters,

3. experimental data,

4. parameters to control the fit and elastic scattering calculations.

Items 1 to 4 may be given in any order. The end of the input for a fit is signaled by a semicolon (;) which causes the fit to begin. When the fit is complete, the potential parameters will be set to the best potential parameters that were found. In addition the predicted optical model scattering cross sections will be printed for each experimental point that was included in the fit. The user may then enter control lines to compute the elastic scattering on a uniform angular grid, or he may increase the accuracy of the calculation (through the use of keywords such as LMINMULT, LMAXADD, STEPSPER or FITACCURACY) and resume the search by entering a second semicolon.

Both the fixed and initial potential parameters are entered using the potential keywords of Section 4.2. The LMINSUB, LMIN, LMAX, LMINMULT, LMAXADD, LMAXMULT, STEPSIZE, STEPSPER and ASYMDTPIA keywords (Section 4.3) may be used to control the accuracy of the elastic scattering calculations during the search. Alternatively one of the elastic PARAMETERSET keywords (Table 4.3) may be used.
5.1. **SPECIFYING THE FIT PARAMETERS AND DATA**

The parameters to be varied in the fit are specified by the FIT keyword. This keyword is followed by a list of potential parameters (Section 4.2) and/or renormalization factors and laboratory angle shifts to be varied. The list must be enclosed in parentheses. If two or more potential parameters are to be held equal to each other during the fit, they should be joined by an equal sign in the FIT list. Some examples of valid FIT specifications are:

- \texttt{FIT (V VI)} - a two parameter fit;
- \texttt{FIT (R0, A=AI, AE=AIE)} - a three-parameter fit with the same energy dependence in A and AI;
- \texttt{FIT (R0=RI0=RC0)} - a one-parameter fit.

Each FIT parameter must have its initial value explicitly entered by means of the potential-defining keywords of Sec. 4.2. The initial value of the first of a string of equal parameters is the one that will be used to start the search.

The normalization of the data and/or the zero point of the laboratory angles may be treated as fit parameters. If only one group of data is being fitted, the keywords RENORMALIZATION and ANGLESIFT may be included in the list of FIT parameters to cause these quantities to be included in the search. (See the description of the DATA keyword below for a precise definition of the meaning of ANGLESIFT.) If there is more than one group of data to be fitted, the keywords RENORMn and/or SHIFTn, where ”n” is an integer from 1 to 20, may be used to indicate which group of data is to be adjusted in magnitude and/or angle. If several groups of data are to be adjusted, then there will be several RENORMn’s or SHIFTn’s in the FIT list; these may be connected with equal signs if the same adjustment is to be made to all of them. The initial values of the renormalizations and shifts that are part of the search are specified in the DATA keyword (see below). Some examples of fits that include searches on the renormalization or angles are:

- \texttt{FIT (V VI A RENORM)} - assumes only one data group.
- \texttt{FIT (R0 R10 SHIFT1=SHIFT2 RENORM1 RENORM2)} - two data groups, both have the same unknown error in angles but possibly different errors in their normalizations.

The default minimizing program (see section 3) cannot be used if the laboratory angle shift is one of the fit parameters. In such cases the POWELL65 or ROCORD minimizer must be specified.

The experimental data are entered using the DATA keyword. This keyword is followed by a pair of parentheses that enclose all of the data that are to be used. As many lines as are necessary may be used to enter the data; the end of the DATA keyword is signaled by the closing parenthesis. If a second DATA keyword occurs in a given job, it will replace, not supplement, the data entered with the first keyword. The data are entered in one or more groups, each containing data for a single elastic channel at a single laboratory energy. Each group is preceded by a list of keywords that give the elastic channel, the laboratory energy, optional overall weight and renormalization factors, and the type of
data to be entered. The data then follow these keywords. The start of the next group is
indicated by the occurrence of a keyword.

The elastic channel is specified by the CHANNEL keyword which is followed by an
elastic channel specification (see section 4.1). The laboratory energy of the group of
data is specified using the ELAB keyword and is in MeV. If there is only one group
of data, the channel and/or laboratory energy may be specified outside (either before
or after) the DATA keyword. Each data item is weighted in the chi-squared sum by
the square of the inverse of its experimental error. An overall weight factor that will
multiply each of these individual weights may be entered using the WEIGHT keyword.
The keyword RENORMALIZATION may be used to enter a renormalization factor that is
multiplied into each experimental value before computing the chi-squared sum. If WEIGHT
or RENORMALIZATION are not entered, the default value of unity is used.

The laboratory angles of the data can be shifted by a constant angle with the ANGLESHPFT
keyword. This keyword specifies an increment (in degrees) that is to be added to each
angle in the laboratory frame. [In the Almagest (ca. 130 AD) PTOLEMY “updated” the
positions of some 1000 stars by shifting the positions of every star by the same amount1.] Input c. m. angles and data are transformed to the laboratory frame for this shift. After
the shift, the Jacobian relating the c. m. and laboratory frames is recomputed at the new
angle, and the angles and data are transformed back to the c. m. frame. Furthermore
if the cross sections are given as ratios to the Rutherford cross sections, the data values
are changed to correspond to the Rutherford cross sections at the new angles. Thus this
keyword treats the data as if the absolute normalization of the data is experimentally
known. If the data was normalized to the Rutherford cross section for small angles, it will
be necessary to use the RENORMALIZATION keyword to specify a suitable renormalization
factor [the average (over the small angles) of the ratios of the Rutherford cross sections
at the original and shifted angles] for the shifted data.

A set of three or four keywords is used to specify the type and order of data being
entered. The set must consist of one keyword from each of the following three groups:

1. ANGLE, CMANGLE, LABANGLE;
2. SIGMA, CMSIGMA, SIGMATORUTH;
3. ERROR, PERCENTERROR, MBERROR.

In addition a fourth keyword, POLARIZATION, may be used to enter polarization data,
but such data will be ignored in the fit. The subsequent data is entered in triples or
quadruples of numbers whose order is the same as that of the three or four keywords.
These keywords remain in effect until a new set is specified; if any one of them is entered,
then a complete new set must be given.

One of the ANGLE, CMANGLE, or LABANGLE keywords is used to indicate the scattering
angles of the data. The ANGLE or CMANGLE keywords are used to designate center-of-mass
angles in degrees. The LABANGLE keyword means that laboratory angles in degrees will
be given. In this case there are two c. m. angles associated with each laboratory angle if

\[ R. \ R. \ Newton, \ The \ Crime \ of \ Claudius \ PTOLEMY, \ (John \ Hopkins \ University \ Press, \ 1977). \]
5.2. CONTROLLING THE FIT

the projectile mass exceeds the target mass. For such cases, positive angles are converted to the smaller possible c. m. angle, while negative angles are converted to the larger. It is not possible to have the cross sections for both c. m. angles added together for comparison with the data.

The type of data that is being entered is indicated by the \texttt{SIGMA}, \texttt{CMSIGMA}, \texttt{LABSIGMA}, or \texttt{SIGMATORUTH} keywords. The first three keywords indicate that cross sections in millibarns are being entered. If \texttt{SIGMA} is used, the cross sections are in the rest frame (c. m. or laboratory) indicated by the \texttt{ANGLE}, \texttt{CMANGLE}, or \texttt{LABANGLE} keywords. The keywords \texttt{CMSIGMA} or \texttt{LABSIGMA} may be used to explicitly indicate the choice of frame or to specify c. m. cross sections at laboratory angles (or vice versa). Ratios of cross sections to the Rutherford (Mott for identical particle scattering) cross sections are indicated by the keyword \texttt{SIGMATORUTH}.

The keywords \texttt{ERROR}, \texttt{MBERROR} or \texttt{PERCENTERROR} are used to indicate the nature of the experimental errors. The \texttt{ERROR} keyword means that errors are being entered in the same units as the data. Thus the errors will be expressed either in millibarns (in the c. m. or laboratory frames as determined by the data) or as a ratio to the Rutherford cross section. The keyword \texttt{MBERROR} indicates that no matter what the data type, the errors are in millibarns. If the data are ratios to Rutherford, then such errors are expressed in the c. m. frame; otherwise the frame is the same as that used for the data. The keyword \texttt{PERCENTERROR} indicates that the errors are expressed in percent.

The following example illustrates the \texttt{DATA} keyword:

\begin{verbatim}
DATA (CHANNEL 16O+40CA ELAB=48
ANGLE PERCENTERROR SIGMATORUTH
 10 5 1.023, 12.5 2 .99 20 3.3 .5
 30 15 .12
ELAB=56 WEIGHT=0.5 ANGLE SIGMA MBERROR
 10 3 .3, 15 1 .1 20 .1 .1 )
\end{verbatim}

Here we are entering data at two different energies. Both sets of data are for $^{16}$O+$^{48}$Ca elastic scattering. The data at the first energy is given at angles of 10, 12.5, 20, and 30 degrees and consists of ratios to the Rutherford cross section that are respectively 1.023, .99, .5, and .12. The errors in these numbers are given as percentages. The data for the second energy are given in millibarns with errors also specified in millibarns. If data that consists of the same quantities in the same order is to be entered at several energies, it is not necessary to repeat the angle, cross section and error keywords for each data group.

5.2 Controlling the Fit

One of the keywords \texttt{LMCHOL}, \texttt{QUAVER}, \texttt{MINIM}, \texttt{DAVIDON}, \texttt{POWELL65}, or \texttt{ROCORD} may be used to pick the minimizing program that is to make the search. The average user will have need of only the default which is \texttt{LMCHOL}. The keywords \texttt{FITMODE}, \texttt{FITMULTIPLE}, \texttt{FITRATION}, \texttt{NUMRANDOM}, and \texttt{REINITIALIZE} are used by some of the following fitters; the default values will almost always suffice. The keywords \texttt{FITACCURACY} and \texttt{MAXFUNCTIONS} are used by all the fitters. A description of these minimizers follows:
CHAPTER 5. OPTICAL MODEL POTENTIAL FITS

1. **LMCHOL** - This is a minimizer that uses analytically computed gradients and makes specific use of the sum-of-squares property of the function that is being minimized. The analytic gradients are computed as the expectation value in the distorted waves of the derivatives of the potential. The potential derivatives are evaluated numerically. **LMCHOL** is based on the Harwell subroutine VA07A which was coded by Fletcher. Despite the fact that the computation of the gradients can more than double the chi-squared sum evaluation time, the **LMCHOL** and **QUAVER** fitters usually find a minimum in less than half the CPU time required by the other fitters. We know of no cases in which the CPU time is significantly longer for **LMCHOL**. The **LMCHOL** fitter is therefore highly recommended and is the default fitter.

2. **QUAVER** - This is a quasi-Newton fitter that uses a pseudo-inverse procedure to solve the required systems of linear equations. The search path followed by **QUAVER** differs significantly from that followed by **LMCHOL** only when there are directions in parameter space for which the second derivative of chi-squared is nearly zero. The keyword **FITRATION** may be used to eliminate steps along such poorly determined linear combinations of the optical model parameters. The default value of $10^{-4}$ will have this effect; smaller values (such as $10^{-15}$) will eliminate such restrictions on the search direction, and result in searches that are almost identical to those of **LMCHOL**. Setting PRINT=2 (section 8) will cause the singular values (the quantities that indicate which linear combinations of parameters are poorly determined) to be printed at each iteration. For both the **QUAVER** and **MINIM** fitters, the keyword **FITMULTIPLE** is a divisor used to reduce the step length when a step to a larger function value is attempted. The default value is 5.

3. **MINIM** - This is a variable-metric fitter using the 1972 Fletcher prescription for the metric update. It is generally slower than the above two fitters.

4. **DAVIDON** - This uses the original variable-metric prescription of Davidon. It is based on the Davidon fitter found in the Argonne Applied Mathematics Division library. This fitter is somewhat slower than the Fletcher fitter (**MINIM**). The keyword **FITMULTIPLE** determines the initial estimate of the metric matrix. The default (**FITMULTIPLE = 0 or FITMULTIPLE > 100**) is to use the second derivative approximation generated from the Jacobian of the chi-squared function. If **FITMULTIPLE** is set to a nonzero value, a diagonal matrix will be used as the initial metric and thus the search will start along the gradient direction. The diagonal elements of the matrix will be $|\text{FITMULTIPLE}| \times D(i)$ where $D(i) = 1$ if **FITMULTIPLE** > 0, and $D(i) = x(i)^2$ if **FITMULTIPLE** < 0 (x designates the parameter vector). The keyword **NUMRANDOM** may be used to specify the number of random steps that are to be made in confirming a minimum. The default is zero.

5. **POWELL65** - This is a minimizer that makes specific use of the sum-of-squares property of the function but does not require PTOLEMY to evaluate the gradient of the function. It is the Harwell subroutine VA02A which is based on a 1965 paper of Powell. The **POWELL65** search algorithm is usually quite efficient for the first
5.3. READING THE OUTPUT

few iterations but then often begins to take very small steps for subsequent iterations. Therefore it often pays to terminate the search intermittently and restart it again. This may be accomplished by using the keyword REINITIALIZE to specify the number of iterations between restarts. REINITIALIZE = 16 is a reasonable value for POWELL65; the default is zero which suppresses reinitialization. The keyword FITMULTIPLE is used to limit the size of a single step; no step will be allowed to exceed FITMULTIPLE×FITACCURACY in relative size. The default is 500, but some tests indicate that larger values (10000) may result in faster searches.

6. ROCORD - This minimizer does not make use of the sum-of-squares property of the function nor of derivatives. It is provided as an alternative should the preceding minimizers fail to behave reasonably. The keyword FITMODE may be used to specify the initial value of IRET for ROCORD; the default is -1021, and should be adequate for most purposes. The keyword NUMRANDOM has the same meaning as for the DAVIDON fitter.

The FITACCURACY keyword may be used to specify the accuracy with which the potential parameters are to be found. Its significance depends upon which minimizer is used but for the first five fitters above it gives the relative accuracy to which each potential parameter or the minimum chi-squared value is to be found. The default value is $10^{-3}$. This keyword is set by the elastic PARAMETERSET’s (section 4.3.). The MAXFUNCTION keyword limits the number of chi-squared sums that may be evaluated during a search. Its default value is 50 which is more than enough for the default fitter (LMCHOL).

5.3 Reading the Output

The output for an optical-model fit begins with a summary of the elastic channels that are being fitted. For each data group, the laboratory energy, the CHANNEL specification, and the type of statistics are listed. The orbital angular momentum range and the stepsize used in solving the Schroedinger equation are also printed. This information is followed by a list of the parameters that are being varied in the fit and their initial values. If data was entered in the laboratory system, the conversion of that data to the c.m. system is shown next. In all subsequent output, the data and fitted values will be printed in the c.m. system only.

One or more pages is then produced for each data group showing the initial values of the fit. The potential parameters evaluated for the channel and laboratory energy are given for each data group. These are followed by a tabulation of the experimental and computed values of the cross sections at each angle involved in the fit. The cross sections are given as ratios to the Rutherford cross section and the angles are c.m. angles. The unsquared contributions to the chi-squared sum are given in a column labeled “(FIT-EXP)/ERROR” the chi-squared sum is the sum of the squares of the entries in this column. The factor WEIGHT is not included in this column. The last column in this tabulation is labeled “AT LMAX” and has been described in Sec. 4.5. Each data group listing is terminated with several lines summarizing the chi-squared for that group. The first
line gives the chi-squared per point (both unweighted and weighted) computed using the RENORMALIZATION entered by the user (the value is also printed in the line). The next line gives that value of RENORMALIZATION that results in the minimum chi-squared (assuming that no other parameters are changed) and the resulting unweighted chi-squared per point. Finally, if there is more than one data group in the fit, the contribution of the current group to the total weighted chi-squared per point is printed, and the total chi-squared per point and chi-squared per degree of freedom for all the data groups are listed at the end of all the data groups.

After this listing of the initial conditions, the course of the fit is summarized by several lines printed at the end of each iteration. The lines give the total chi-squared per point and the corresponding parameter values (the parameters are listed in the order they were specified in the FIT keyword – this order may be found in the top right-side of the header on each page). If the listing of iterations terminates with the message

FIT COMPLETE. FINAL RESULTS FOLLOW

then the minimizer being used located a minimum of chi-squared and terminated normally. Any other message indicates an abnormal termination. The most likely cause of an abnormal termination is the failure of the minimizer to locate a minimum within the allowed number of function references (steps); such a condition is indicated by the message

NOT ENOUGH ITERATIONS TO ACHIEVE A GOOD FIT

In such cases the parameter values at the time of termination of the minimizer are used as the final parameter values; these will not be parameters corresponding to a minimum of $\chi^2$. PTOLEMY then gives a listing of the final parameter values and of the resulting fit; the format of this listing is identical to that used for the initial conditions.

A page labeled FINAL VALUES AND UNCERTAINTIES follows the listing of the fit. This page has a column for each fit parameter; the columns are labeled with the parameter names. The first line of the table contains the final value of each parameter. Next are several lines giving the gradient at the minimum; the gradient is ideally zero but if there are large second derivatives may be significantly non-zero. Then there are two lines giving the RMS uncertainty and the relative RMS uncertainty in the parameters. These are based on the diagonal elements of the covariance matrix (error matrix) and are defined as the changes in the parameter values that would result in the $\chi^2$ (not $\chi^2$ per point) increasing by unity. Finally the eigenvalues and eigenvectors of the relative covariance matrix are listed. Each row of this listing contains the square root of an eigenvalue and the corresponding eigenvector. The square root of the eigenvalue is the relative change in the linear combination of parameters specified by the eigenvector that would result in the $\chi^2$ increasing by unity. Thus eigenvectors with small eigenvalues represent well-determined combinations of parameters.
Chapter 6

Collective Model DWBA for Inelastic Excitation

PTOLEMY is capable of doing DWBA calculations for inelastic-scattering reactions. The collective-model form factor is used for the nuclear part of the effective interaction. Only one nucleus may be excited in the reaction and that nucleus must initially have spin zero. The multipolarity (referred to as $L_x$) of the excitation must be greater than zero, but is otherwise not limited. The effective interaction that causes the excitation is the radial derivative of the Coulomb and both real and imaginary nuclear parts of an optical potential. Usually this optical potential will be that of the incoming channel, but a different potential may be used.

6.1 Specifying the Physical Problem

In a standard PTOLEMY calculation of an inelastic-scattering reaction, the input will be in the following order:

1. Masses, charges, etc., of the 4 nuclei.
2. Grid and other calculation-control parameters.
3. Optical potentials for the two scattering states.
4. Optional potential changes for effective interaction.

Variations of this order are possible: item 4 is not required and item 2 may be spread among the other items.

In more detail a typical PTOLEMY input deck for inelastic excitation will look like

REACTION: reaction definition, ELAB = ...
PARAMETERSET...., other parameter specifications
INCOMING incoming optical parameters ;
Here the keywords INCOMING, and OUTGOING indicate which potential parameters are being entered. The semicolons indicate that the complete potential has been defined and that PTOLEMY is to go ahead with that stage of the computation (the semicolons are actually part of the input). The final semicolon indicates that all of the computational parameters have been entered and that the DWBA calculation should begin. Of course each of the potential specifications will in general require more than one input line.

The easiest way to define the participants in the reaction is to use the REACTION keyword. This keyword is followed by the statement of the reaction in standard nuclear terminology. Some examples will best illustrate the possibilities:

\textbf{REACTION}: \texttt{48CA(C12, 12C)CA48(2+ 3.83)}

\textbf{REACTION} = \texttt{208PB(16O, 16O(6.13, 3-))208PB}

All four nuclei must be included in the REACTION specification, and the complete reaction specification must be on the same input line as the REACTION keyword. Excited state information is enclosed in parentheses with no spaces or other punctuation separating the nuclear symbol and the left parenthesis. Except as has been otherwise indicated, blank spaces and commas may be freely used to make the reaction specification more readable. See the description of the CHANNEL keyword (section 4.1) for details concerning the nuclide symbols and excited-state specifications.

The REACTION keyword results in the definition of the atomic mass, charge, spin, and excitation energy of the four particles. Individual data values defined by the REACTION keyword may be overridden by the use of other keywords or the REACTION keyword may be omitted and all of the particle definitions entered via other keywords. These keywords are described in the chapter on transfer reactions (Section 7.1).

The scattering energy is entered by either of the keywords ELAB or ECM followed by the energy in MeV. In both cases the energy refers to the incident kinetic energy; the outgoing energy is determined from the $Q$ value of the reaction. The excitation energy is normally used by PTOLEMY to determine the $Q$ value of the reaction. If desired, a different $Q$ value may be entered by using the keyword $Q$ followed by the value in MeV.

The optical potentials for incoming and outgoing states are entered using the potential defining keywords of Sec. 4.2. The keywords INCOMING and OUTGOING indicate which state is being defined and a semicolon (;) is used to indicate the end of a particular definition. There are no default values for the radius or diffuseness parameters. The well depths are all zero by default. The potential parameters used for the incoming state will also be used for the outgoing state unless they are explicitly overridden. While this is generally desired for heavy-ion reactions, for which PTOLEMY was written, it can have unexpected consequences in light-ion reactions. The laboratory energy corresponding to the outgoing scattering energy is used for the outgoing scattering parameters so that the same keyword
values will give slightly different potentials in the incoming and outgoing channels if an energy dependence is specified. As an example, the input lines

\[
\text{INCOMING V=50 VI=25 R0=1.2 A=.5 RCO=1.2 ;}
\]

\[
\text{OUTGOING ;}
\]

define the same optical potential for both the incoming and outgoing states. Since the radius and diffuseness of the imaginary potential are not given, they are chosen to be the same as those of the real potential.

The effective interaction for the inelastic excitation is:

\[
H(r) = -\beta \left[ R' \frac{d(V_{\text{real}})}{dr} + R'_I \times \frac{d(V_{\text{imag}})}{dr} \right] + \beta C R'_C 3Z A \frac{Z Ae^2}{(2L_x + 1)f(r)}
\]

where

\[
f(r) = \begin{cases} 
\frac{r^{L_x}}{R_C^{(L_x+2)}} & r < R_C, \\
\frac{R_C^{(L_x-1)}}{r^{(L_x+1)}} & r \geq R_C,
\end{cases}
\]

Here \( R' \) is the nuclear radius of the nucleus that is being excited:

\[
R' = R_0 \times A'^{(1/3)}
\]

\[
R'_I = R_{I0} \times A'^{(1/3)}
\]

\[
R'_C = R_{C0} \times A'^{(1/3)}
\]

(6.3)

where the \( R_0 \) are the radius parameters used in the effective interaction. The atomic weight of the excited nucleus is denoted by \( A'(A' = M_a \text{ or } A' = M_A) \). The quantities \( \beta \times R', \beta \times R'_I, \text{ and } \beta C \times R'_C \) are the so-called deformation lengths of the excited nucleus. The definition of \( f(r) \) is that obtained from a multipole expansion of a point charge interacting with a uniform sphere of charge. Note that the full RC (not \( R'_C \)) appears in \( f(r) \).

The dimensionless Coulomb and nuclear deformation parameters may be entered with the keywords BETAC and BETA respectively. Alternatively the Coulomb deformation may be specified by the keyword BELX which enters the \( B(E, L_x) \), in units of \( e^2\text{barn}L_x \), for the corresponding electromagnetic excitation process:

\[
\text{BELX} = B(E, \text{ excitation}) = \frac{[2J(\text{final}) + 1]}{[2J(\text{initial}) + 1]} B(E, \text{ decay})
\]

(6.4)

The value of BETAC is then computed from

\[
\beta_c = \frac{4\pi}{3Z} \times \sqrt{\text{BELX}} \times \left( \frac{10}{R'_C} \right)^{L_x} \sqrt{\frac{[2J(\text{initial}) + 1][2L_x + 1]}{[2J(\text{final}) + 1]}}
\]

(6.5)
where $Z$ is the atomic number of the excited nucleus and $R'_C$ (in fm) is defined above. If both BETA and BETAC (or BELX) are not specified, then the missing parameter is chosen such that the deformation lengths are equal:

$$\beta R' = \beta C R'_C$$

(6.6)

where $R'$ is the nuclear radius of the excited nucleus.

The optical potential that appears in the effective interaction is the optical potential that was used in the incoming state. It may be modified by entering different potential parameters after the semicolon that defines the outgoing state. If a potential parameter is not so changed, its value in the incoming state will be used. Thus

\begin{verbatim}
INCOMING V=50 VI=20 R0=1.2 A=.65 RC0=1.2 ;
OUTGOING ;
VI = 1E-6 ;
\end{verbatim}

will result in the effective interaction being essentially real. (Note that it is unfortunately not possible to redefine the well depths to be exactly zero.)

### 6.2 Controlling the Calculation

The keywords LMIN and LMAX, or MINMULT, LMINSUB, LMAXMULT, and LMAXADD determine the range of $L$-values for which both the nuclear and Coulomb excitation amplitudes are computed. These keywords are explained in section 4.3, and reasonable values may be found in Table 6.2 below. The $L_C$ used in the formulas in section 4.3, is the average of the critical $L$-values in the incoming and outgoing channels.

The Coulomb excitation amplitudes generally must be found for much larger values of $L$ than the LMAX used for the nuclear amplitudes. Using a semi-classical approximation\(^1\), ptolemy estimates an $L'_{\text{max}}$ such that the Coulomb amplitudes are negligible. The keyword DWCUTOFF may be used to control this choice; the choice is made such that

$$|\text{Amplitude}(L'_{\text{max}})| = \text{DWCUTOFF} \times |\text{Amplitude}(L_c)|,$$

(6.7)

where both amplitudes are the pure Coulomb amplitudes. Pure Coulomb excitation amplitudes (evaluated using Coulomb scattering wavefunctions) are used for $\text{LMAX} < L < L'_{\text{MAX}}$.

For $\text{LMIN} \leq L \leq \text{LMAX}$, the nuclear and Coulomb amplitudes are evaluated as a one-dimensional integral over the range $\text{SUMMIN} < R < \text{SUMMAX}$. The Coulomb contribution for $\text{SUMMAX} \leq R < \infty$ is then evaluated using the asymptotic expansion described later. The lower limit ($\text{SUMMIN}$) is usually picked as that value of the radial coordinate beyond which the elastic scattering wavefunction for $\text{LMIN}$ exceeds $10^{-15}$ in magnitude. The

---

\(^1\)K. Alder, A. Bohr, T. Huss, B. Mottelson and A. Winther, Rev. Mod. Phys. 28, 432 (1956), Eq. 2.83
upper limit (SUMMAX) is chosen to be the value of ASYMPTOPIA specified for the scattering wavefunctions. These two choices may be overridden with the keywords SUMMIN and SUMMAX, but there should be no need to do so. The number of Gauss points used in the one-dimensional integral is specified by the keyword SUMPOINTS which gives the number of points to use per average wavelength in the incoming and outgoing channels. SUMPOINTS does not need to be an integer. The Gauss points are mapped into the interval (SUMMIN, SUMMAX) using mappings defined in section 7.2 which are controlled by the keywords MAPSUM, GAMMASUM, and SUMMID. The default value of SUMMID is the midpoint of the (SUMMIN, SUMMAX) interval. There should never be any need to override the default mapping procedure.

The computation outlined in the previous paragraph leaves the following Coulomb excitation integrals to be evaluated:

1. \( L_{\text{min}} \leq L \leq L_{\text{max}}, \text{SUMMAX} \leq r < \infty \);
2. \( L_{\text{max}} < L \leq L'_{\text{max}}, 0 \leq r < \infty \).

In case 1) it is necessary to do these integrals for both the regular and irregular Coulomb wavefunctions; linear combinations, based on the elastic incoming and outgoing S-matrix elements, of the resulting amplitudes are then used. For case 2) we assume that the scattering is determined by the point Coulomb charges alone and only the regular Coulomb wavefunctions are used.

In both cases the required Coulomb excitation amplitudes are found by recursion relations on \( L \). These recursion relations are quite fast and are reasonably stable. The starting values of the recursion relations are found by a combination of numerical integrations and an asymptotic expansion\(^2\) for integrals of Coulomb wavefunctions and inverse powers of \( r \). The numerical integrals are done cycle by cycle until the asymptotic expansion may be used; the number (which must be an integer) of Gauss points used in each wavelength is entered using the keyword NPCOULOMB. The relative accuracy required of the asymptotic expansion may be controlled by the keyword INELASACC. In addition a test of the recursion relations is made by explicitly computing the final recursed values. If the recursion relation is in error by more than \( 10 \times \text{INELASACC} \), a warning message is printed. Reasonable values of NPCOULOMB and INELASACC may be found in Table 6.2 below.

The computation of the scattering wavefunctions may be controlled with the ASYMPTOPIA and STEPSIZE or STEPSPER keywords. These keywords are defined in section 4.3. As has just been described, the nuclear part of the excitation amplitude is integrated out to ASYMPTOPIA; by the use of the asymptotic Belling expansion, the Coulomb excitation amplitude is integrated to infinity.

The keyword PARAMETERSET (Section 4.3) may be used to select standard groups of grid-setting parameters. Table 6.2 gives the PARAMETERSET names and the associated values that are suitable for inelastic excitation calculations.

Since the PARAMETERSET groups of Table 6.2 define values of ASYMPTOPIA and STEPSPER that are to be used in the two-body states, the PARAMETERSET keyword should precede the definitions of the two-body states.

---

\(^2\) J.A. Belling, J. Phys. B 1, 136 (1968)
CHAPTER 6. COLLECTIVE MODEL DWBA FOR INELASTIC EXCITATION

<table>
<thead>
<tr>
<th>Keyword</th>
<th>Default</th>
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<th>INELOCA2</th>
<th>INELOCA3</th>
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<tr>
<td>STEPSPER</td>
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<td>25</td>
</tr>
</tbody>
</table>

Table 6.1: **PARAMETERSET** names and associated values for inelastic scattering calculations. The first column gives the default values.

6.3 Reading the Output

As each of the two scattering states are entered, a summary of the two-body channel and the potential is printed; this summary was described in Sec. 4.5. The computation of the inelastic excitation amplitudes is preceded by a page summarizing the reaction. This page contains a listing of the nuclei involved in the reaction, the deformation parameters, and the potential parameters used for the effective interaction. The column labeled DEPTH in the latter contains $-3Z_aZ_Ae^2$ for the Coulomb part of the effective interaction. Next the range of angular momenta (LMIN, LMAX) for which both the nuclear and Coulomb amplitudes are computed is listed. This is followed with a summary of the one-dimensional integration grid that is used to compute the non-asymptotic part of these amplitudes.

Next a summary of the determination of the maximum L value ($L'_{MAX}$) needed for the Coulomb amplitudes is given. The maximum value required for each ($L_x, L_{out} - L_i$) pair is given; the maximum of all of these values is then used. Following this summary one or more warning messages of the form

**FOR LIN, LOUT = RECURSION IS POOR:**

may be printed. These indicate that the recursed values of the Coulomb amplitudes did not compare well with the explicitly computed values. Both values are printed in the message. If the difference of the two values is not large, or if the values are both unusually small, then the message may be ignored. Otherwise the calculation should be repeated using a larger value of NPCOULOMB and/or a smaller value of INELASACC. A page labeled INTERPOLATION AND EXTRAPOLATION IN L contains the line MAXIMUM L0 USED IN COMPUTING.... This line gives the maximum value of L ($L'_{MAX}$) for which the Coulomb amplitudes were computed by the recursion relations.

The pages labeled REACTION AND ELASTIC PARTIAL WAVE AMPLITUDES give the amplitudes for LMIN ≤ L ≤ LMAX. ****NEEDS WORK**** The columns labeled RADIAL INTEGRAL give the magnitude and phase (in radians) of the inelastic excitation amplitude. This is the amplitude defined in Eq. 2.39 except that a factor ($\beta + \beta_{AC}$)/2 has
been removed. The columns labeled **INCOMING ELASTIC** and **OUTGOING ELASTIC** give the magnitudes and phase-shifts (in radians) of the elastic S-matrix elements. The Coulomb phase shifts are also given in radians.

The last set of pages give the inelastic excitation cross sections. These are given (in millibarns) in a column labeled **REACTION** and are c. m. values. The column labeled **LOW L %/L** has the significance described in Sec. 4.5, however the column labeled **% FROM L > LMAX** has a quite different meaning from the corresponding column in the elastic scattering output. Here the column gives (as a percentage) the total contribution of the (pure Coulomb) amplitudes for **LMAX < L ≤ LMAX’**. It is not to be construed as an indication of error. The columns labeled **INCOMING/RUTHERFORD** and **OUTGOING/RUTHERFORD** give the elastic cross sections relative to the Rutherford values.

Following the tabulation of the differential cross sections is a line labeled **TOTAL**. This line gives the total inelastic excitation cross section (computed by summing the partial-wave amplitudes – not by integrating the printed angular distribution) and the total reaction cross sections for the entrance and exit channels. Following this line is a breakdown of the total excitation cross section into contributions from each magnetic substate (the axis of quantization is the incoming beam direction). Although values are listed only for **Mx ≥ 0**, the listed values for **Mx > 0** are not doubled.
Chapter 7

Finite Range DWBA for Transfer Reactions

PTOLEMY can carry out finite range DWBA calculations for particle transfer reactions. The reaction may be either stripping or pickup, and may involve more than one exchanged orbital angular momentum (referred to as $L_x$). Either the post or prior approximation may be used, and the effective interaction may include terms for the Coulomb part of the bound-state potential, the core-core Coulomb optical potential, and the real part of the core-core nuclear optical potential. When all these terms are included, the discrepancy between post and prior calculations is usually less than a few percent. Core-core terms for the imaginary part of the optical potential may not be included.

7.1 Specifying the Physical Problem

In a standard PTOLEMY calculation of transfer reactions, the input will be in the following order:

1. Masses, charges, etc., of the 5 particles.
2. Integration grid specifications and $L$-value ranges.
3. Potentials for the two bound states.
4. Optical potentials for the two scattering states.

Variations on this order are possible; the most likely is the spreading of item 2) among the other items.

In more detail a typical PTOLEMY input deck will look like

```plaintext
REACTION: reaction definition, ELAB = ...
PARAMETERSET ...., other computation parameters
PROJECTILE projectile bound state parameters ;
```
7.1. **SPECIFYING THE PHYSICAL PROBLEM**

TARGET target bound state parameters;

INCOMING incoming optical parameters;

OUTGOING outgoing optical parameters;

. ;

RETURN

Here the keywords PROJECTILE, TARGET, INCOMING, and OUTGOING indicate which potential parameters are being entered. The semicolons indicate that the complete potential has been defined and that PTOLEMY is to go ahead with that stage of the computation (the semicolons are actually part of the input). The final semicolon indicates that all of the computational parameters have been entered and that the DWBA calculation should begin. Of course each of the potential specifications will in general require more than one input line.

The easiest way to define the participants in the reaction is to use the **REACTION** keyword. This keyword is followed by the statement of the reaction in standard nuclear terminology. Some examples will best illustrate the possibilities:

**REACTION:** 48CA(160, 14C)50TI

**REACTION =** 208PB(016,15N)BI209(7/2- .90)

**REACTION PB208(016 15N(3/2, 6.3239))BI209**

**REACTION PB208(16O(2,6.93) 12C)PO212**

All four nuclei must be included in the **REACTION** specification, and the complete reaction specification must be on the same input line as the **REACTION** keyword. Excited state information is enclosed in parentheses with no spaces or other punctuation separating the nuclear symbol and the left parenthesis. An excited state of the initial target may not be given in this manner, but any or all of the other three particles may have excited state descriptors. See the description of the **CHANNEL** keyword (Section 4.1) for details concerning the nuclide symbols and excited-state specifications. Except as has been otherwise indicated, blank spaces and commas may be freely used to make the reaction specification more readable.

The **REACTION** keyword results in the definition of the atomic mass and charge of the four particles. The atomic mass and charge of the exchanged particle is then computed by subtraction. The **1975 Oak Ridge Atomic Mass Adjustment** and the 1971 Nuclear Wallet Cards compilation\(^1\) is then used to find the ground state mass excesses and spins of all 5 particles. The ground state mass excesses (along with the excitation energies, if given) will be used during the bound state computation to find the separation energies of the exchanged particle (X) from the appropriate cores.

Individual data values defined by the REACTION keyword may be overridden by the use of other keywords or the REACTION keyword may be omitted and all of the particle definitions entered via other keywords. The keywords that define the five particles have the form "Ki" where "K" indicates what quantity is being defined and "i" is a suffix indicating which particle is involved (A, A, B, B, or X). The possibilities for K are

- M - The mass in AMU. This need not be an integer.
- Z - The charge.
- J - The spin of the nucleus.
- MXCG - The mass excess of the ground state in MeV.
- E* - The excitation energy in MeV.
- MXC - The mass excess of the nucleus in MeV (MXCi = MXCGi + E*i).

As an example MB=15, ZB=7, JB=3/2, E*B=6.3239, MXCGB = .10152 would define the excited state of 15N contained in the third REACTION example given above. (Alternatively one could have entered MXCB = 6.42542 and left out the E*B and MXCGB keywords or one could directly enter the proton separation energy at the time of the 16O bound state calculation and leave out all the mass excess specifications.) In the following example

REACTION: 209Bi(16O 12C)213At, JBIGB = 9/2

one is supplying the ground state spin of 213At which is not in PTOLEMY’s table.

The scattering energy is entered by either of the keywords ELAB or ECM followed by the energy in MeV. In both cases the energy refers to the incident kinetic energy; the outgoing energy is determined from the Q value of the reaction.

The Q value may be entered by using the keyword Q followed by the value in MeV. If it is not entered it will be found as the difference of the two bound state energies. If it is entered then it is necessary to define only one of the bound state energies; the other will be found using the Q value. Since the REACTION keyword results in the definition of both bound state energies, it is usually not necessary to enter the Q value.

The version of the DWBA on which PTOLEMY is based uses as the effective interaction that induces transfer the potential that binds the composite particle at either the projectile or target vertex. The vertex whose potential is to be used as the effective interaction is specified by the keywords USEPROJECTILE or USETARGET. USEPROJECTILE indicates that the potential for the projectile bound state is the interaction potential; USETARGET causes the target potential to be used. The default is USEPROJECTILE. Note that we avoid the use of the words “post” and “prior” in specifying the interaction vertex.

The content of the interaction potential is controlled with the NUONLY, USESIMPCOULOMB, USECOULOMB or USECORE keywords. NUONLY means that only the nuclear part of the bound state potential is used in the interaction potential. (Most published DWBA calculations have used this prescription.) USESIMPCOULOMB means that the full bound state potential at the vertex designated by USEPROJECTILE or USETARGET is used. USECOULOMB causes the nuclear potential at the designated vertex to be used with the complete three-body Coulomb potential. The inclusion of the Coulomb corrections due to the third
7.1. SPECIFYING THE PHYSICAL PROBLEM

particle removes post/prior discrepancies from the Coulomb part of the interaction and can result in much closer agreement between USETARGET and USEPROJECTILE results. The keyword USECORE may be used to include both the Coulomb and core-Coulomb corrections that are included by the USECOULOMB keyword and the core corrections from the real part of the nuclear optical potential. Core corrections due to the imaginary part of the optical potential (which are typically only a few percent) cannot be included. The default is USECORE. Note that DWBA calculations are usually somewhat simpler numerically if the interaction is attached to the vertex (usually the projectile vertex) involving the lighter ions. In a reaction such as $^{208}$Pb($^{16}$O, $^{17}$O)$^{207}$Pb in which the projectile and target differ significantly in mass, a USETARGET calculation requires a substantially denser integration grid than is required for USEPROJECTILE, however if the USECORE option is selected, essentially the same cross sections will be found in both cases.

The spectroscopic amplitudes for the two bound states may be entered with the keywords SPAMP and SPAMT. These amplitudes will be squared and multiplied into the cross sections. The default values are unity. Alternatively the spectroscopic factors (the squares of the amplitudes) may be directly entered using the keywords SPFACP and SPFACT. The spectroscopic factors must be positive but the amplitudes may be negative.

Potential parameters must be entered for each of the two bound states and the two scattering states. The potentials are defined when the bound state or scattering state is to be computed; the same keywords are reused to define the potentials in each of the four states. Each of these four two-body states consists of two particles that are referred to as the “projectile” and “target” (not to be confused with the projectile bound state and target bound state). In the scattering states these words have their normal meanings; for the bound states the exchanged particle (X) is always the “projectile”. The potential parameters are defined in Sec. 4.2. The laboratory energy corresponding to the outgoing scattering energy is used for the outgoing potential parameters so that the same keyword values will give slightly different potentials in the incoming and outgoing channels if an energy dependence is specified.

If a given $V$ or $\tau$ is defined in a channel, then its associated $R$ (or $R_0$) and $A$ must also be defined. At the beginning of input for each channel, all $V$’s and $\tau$’s are set to 0 and all $R$’s and $A$’s are undefined. However, potential parameters will be retained (if they are not overridden) from one bound state to the other and from one scattering state to the other. They will not be retained from bound states to scattering states or vice-versa. The same rules apply to $RC$ and $RC_0$, one of which must be defined for the bound states if both $Z_p$ and $Z_t$ are nonzero. If both $RC$ and $RC_0$ are undefined for the scattering states, the Coulomb potential of two uniform spheres (Section 6b) will be used.

Normally the binding energy (cluster separation energy) of the bound states will be computed by PTOLEMY from the information in the REACTION specification. If it is desired to override this bound state energy, one may use the $E$ keyword to enter the bound state energy along with the bound state potential. The energy is given in MeV and must be negative for bound states. The $E$ keyword may also be used in the scattering state descriptions in which case it specifies the c. m. energy of the state. It will then override the c. m. energy determined from the ELAB keyword or from ELAB combined with the $Q$ value depending on the channel. One may use the $E*P$ or $E*T$ keywords to enter the
excitation energy of the projectile or target. Use of FITE (page 38.) in DWBA calculations may result in bound state energies (and hence $Q$-values) that are significantly different from the values determined from the REACTION specification. The default is FITV.

### 7.2 The Integration Grid

The PTOLEMY integration grid is constructed by a rather elaborate set of subroutines consisting of some 1100 FORTRAN source cards. The construction is fairly automatic and is designed to place the integration points where the integrand is largest. Ideally the user would not have to intervene in this process and would only have to tell the processor the desired accuracy of the final results (the differential cross sections). Unfortunately the method is less than perfect and the user must have some understanding of what is done. This section describes most of the parameters that control the grid construction. The novice user of PTOLEMY need not be concerned with the details of this section since the PARAMETERSET keyword described in the following section can be used to choose complete sets of grid-construction parameters. As is described in the following section, these parameters have been tuned for specific types of calculations; they may not be adequate for quite different types of calculations, and the convergence of the computed cross sections should be checked by repeating such calculations with a different set of parameters.

The PTOLEMY integration grid is based on the three variables $d$ (DIF), $s$ (SUM), and $\phi$ (PHI):

\[
\begin{align*}
    d &= R_i - R_{out} \\
    s &= (R_i + R_{out})/2 \\
    \phi &= \text{angle between } R_i \text{ and } R_{out}.
\end{align*}
\]

Here $R_i$ and $R_{out}$ are the radial variables in the incoming and outgoing elastic channels. The form factor is first integrated over $\phi$. This results in a quantity that, for fixed values of $d$, is a smoothly varying function of $s$. However the scattering wavefunctions are often rapidly varying functions of $s$. Therefore the form factor is computed on a rather coarse grid of $s$ values and interpolated to a fine grid for integration with the scattering wavefunctions. The numbers of points in each of the grids on which the form factor is computed are specified by the keywords NPSUM, NPDIFF, and NPPHI, respectively. These numbers may have any value between 1 and 2000; typical values will be found in Table 7.3 below. The number of points in the SUM integration grid is specified by the keyword SUMPOINTS which gives the number of points per average wavelength in the entrance and exit channel. In this manner, the grid automatically becomes denser as the bombarding energy is increased; it is our experience that the other three grids (controlled by NPDIFF, NPSUM and NPPHI) need not become denser as the bombarding energy increases. The same three-dimensional grid is used for all values of $L_i$, $L_{out}$ and $L_x$.

The keyword DWCUTOFF is used in the construction of the integration grids. DWCUTOFF specifies in a relative sense the smallest integrand

\[
(R_i \times \text{scattering wave} \times \text{bound state} \times \text{potential} \times \text{bound state} \times \text{scattering wave} \times R_{out})
\]
to include in the grid. If the integrand at a point \((R_i, R_{out}, \phi)\) is smaller than \(\text{DWCUTOFF}\) times the largest value of the integrand encountered, that point will not be included in the grid.

The lower and upper limits of the \(\text{SUM}\) grid are determined as the values of \(s\) for which the integrand has fallen (in a relative sense) beneath \(\text{DWCUTOFF}\) at \(d = \phi = 0\). The scattering wavefunctions for \(L = \text{LMIN}\) are used when finding the lower limit of \(s\) while the wavefunctions for \(L = L_c\) are used for the upper limit. These two limits may be overridden by using the \(\text{SUMMIN}\) and/or \(\text{SUMMAX}\) keywords to give values (in fm) of the lower and upper limits.

The \(\text{SUM}\) grid points are mapped into the interval \((\text{SUMMIN}, \text{SUMMAX})\) in a manner that clusters them about a “midpoint.” This midpoint is at present chosen to be the expectation value of \(\text{SUM}\) (weighted by the integrand for \(L = L_c\)) for \(d = \phi = 0\). The value of the “midpoint” may be overridden with the keyword \(\text{SUMMID}\) or it may be multiplied by a factor specified by the \(\text{MIDMULT}\) keyword, however the resulting value will always be constrained to be not greater than the average of \(\text{SUMMIN}\) and \(\text{SUMMAX}\).

The form of mapping used for the \(\text{SUM}\) grid may be controlled by the \(\text{MAPSUM}\) keyword which may have one of the following values:

- 0 - linear mapping with no compression.
- 1 - cubic mapping with sinh compression.
- 2 - rational mapping with sinh compression (default).
- 3 - linear mapping with sinh compression.

\(\text{SUMMAP} = 2\) gives the best results in the cases so far examined and is the default. The degree of compression in mappings 1-3 is controlled by the \(\text{GAMMASUM}\) keyword. Suitable values are indicated in Table 7.3 below.

The \(\text{DIF}\) grid limits are determined by the values of \(d\) for which the form factor becomes smaller in a relative sense than \(\text{DWCUTOFF}\). Initially these limits are chosen separately for each of the \(\text{NPSUM}\) values of \(S\). Each \(\text{DIF}\) grid is also mapped about a ”midpoint” which is chosen to be the location of the maximum of the form factor for the fixed value of \(\text{SUM}\) (and \(\phi = 0\)). Since the automatic choice of the \(\text{DIF}\) limits and midpoints seems always to be successful, no keywords are provided to override these values. The keyword \(\text{MAPDIF}\) is used to choose the \(\text{DIF}\) mapping and has the same meanings as the \(\text{MAPSUM}\) keyword. The best value in the cases studied is 1 which is the default. The keyword \(\text{GAMMADIF}\) specifies the degree of compression used in the mapping.

To allow interpolation in the \(\text{SUM}\) variable, the \(\text{DIF}\) grid points must be smoothly varying functions of \(S\). The procedure outlined in the previous paragraph does not necessarily produce such a grid. Therefore low-order polynomials in \(S\) are fitted to the families of \(\text{DIF}\) points that are chosen above. These polynomials are then used to generate the \(\text{DIF}\) points that are actually used in the calculation. The order of these interpolating polynomials may be specified by the \(\text{VPOLYORDER}\) keyword; the default is 3 which is almost always adequate. (The error message

\text{INVALID VMIN, VMID, VMAX...}
usually means that \texttt{VPOLYORDER} should be increased, although the cross section values are often only slightly affected by this error.)

The \( \phi \) maps are individually chosen for each value of \( s \) and \( d \). The minimum \( \phi \) is always 0 and the maximum is determined as the point at which the form factor falls in a relative sense beneath \texttt{DWCUTTOFF}. For heavy ion reactions this is usually a small angle \( [\cos(\phi) > .99] \). The \( \phi \) map is a linear map in the variable \( \cos(\phi) \).

### 7.3 Controlling the Calculation

The keywords \texttt{LMIN} and \texttt{LMAX}, or \texttt{LMINMULT}, \texttt{LMINSUB}, \texttt{LMAXMULT}, and \texttt{LMAXADD} determine the range of \( L \)-values for which the transfer amplitudes are computed. These keywords are explained in section 4.3, and reasonable values may be found in Table 7.3 below. The \( L_{\text{critical}} \) used in the formulas in section 4.3 is the average of the critical \( L \)-values in the incoming and outgoing channels.

Not all radial integrals used in computing the differential cross sections need be explicitly computed by \texttt{PTOLEMY}. \texttt{PTOLEMY} will interpolate between computed values and extrapolate beyond the largest value of \( L \) for which the radial integrals are computed. The keyword \texttt{LSTEP} determines which radial integrals are to be computed. The radial integrals for

\[
L_{\text{OUT}} = \text{LMIN}, \text{LMIN + LSTEP}, \text{LMIN + 2LSTEP}..., \quad \text{(where the sequence stops at or before } L_{\text{MAX}} \text{)}
\]

will be computed. The radial integrals for all \( L_x \) and \( L_i \) associated with these \( L_{\text{OUT}} \)'s will be computed. The remaining radial integrals for \( \text{LMIN} \leq L_{\text{OUT}} \leq \text{LMAX} \) will then be found by interpolation using continued fractions. The default value of \texttt{LSTEP} is 1 which causes all radial integrals from \texttt{LMIN} to \texttt{LMAX} to be computed explicitly.

In addition \texttt{PTOLEMY} will pick a \( L'_{\text{MAX}} > \text{LMAX} \) such that radial integrals for \( L_i, L_{\text{OUT}} > L'_{\text{MAX}} \) are negligible. The radial integrals for \( \text{LMAX < L_i, LOUT \leq L'_{MAX}} \) will be found by extrapolation. The extrapolating function used is of Woods-Saxon form in \( L \). Thus \texttt{LMAX} must be sufficiently beyond the \( L \)-window for such a shape to be an adequate representation of the radial integrals. The keyword \texttt{MAXEXTRAPOLATION} may be used to limit or completely suppress the extrapolation to \( L > \text{LMAX} \). It specifies the maximum allowed \( L'_{\text{MAX}} - \text{LMAX} \). If it is set to 0, no extrapolation will occur. The default value is 100.

Since no extrapolation to \( L_i, L_{\text{OUT}} < \text{LMIN} \) is made, \texttt{LMIN} must be small enough to include all important radial integrals. Care should be taken to avoid specifying too small a value of \texttt{LMIN} since the radial integrals for small \( L_i \) and \( L_{\text{OUT}} \) are small due to extensive cancellations of the integrand viewed as a function of \( R_i + R_{\text{OUT}} \). In practice these cancellations are hard to reproduce without using a large number of Gauss points and the computed radial integrals may be much larger than they should be. Thus a more accurate solution is often obtained by totally excluding integrals whose contributions are very small but which are hard to calculate accurately.

For a given reaction there will usually be several possible values of the transferred orbital angular momentum:

\[
L_x = L_{\text{OUT}} - L_i = J(\text{projectile}) - J(\text{target})
\]
PTOLEMY will compute the radial integrals for all possible values of $L_x$ and add the resulting cross sections together with the appropriate Racah coefficients. If only one value of $L_x$ is possible the resulting cross section will still be weighted by the Racah coefficient. If it is desired to have results for only one value of $L_x$, the keyword $LX$ may be used to specify the desired value.

The computation of the two-body wave functions (both bound and scattering states) may be controlled with the ASYMP_TOPIA and STEPSIZE or STEPSPER keywords. These keywords are defined in section 4.3. ASYMP_TOPIA and STEPSIZE or STEPSPER may be respecified for each of the four two-particle states; if they are not reentered, the value last entered is used.

The value of ASYMP_TOPIA in effect when the bound states are specified is the largest value of $R_p$ or $R_t$ (the radial coordinates of the projectile and target bound-state wavefunctions) at which the bound-wave functions will be found and thus must be large enough to satisfy the needs of the integration grid. The largest values of $R_p$ and $R_t$ used by the integration grid are printed in the summary of the grid.

The largest value of $R_i$ and $R_{out}$ (the scattering variables) will automatically be chosen to be large enough to satisfy the needs of the integration grid. However it will never be smaller than the value of ASYMP_TOPIA in effect at the time of input of the optical potentials. Therefore since, as has just been explained, a large ASYMP_TOPIA is often needed for the bound-state wave functions, substantial memory savings may be realized by reducing ASYMP_TOPIA to as small a value as is physically reasonable for the scattering states; it will then be automatically increased to the required value.

The keyword PARAMETERSET (Section 4.3) may be used to select standard groups of grid setting parameters. Table 7.3 gives the PARAMETERSET names and the associated values that are suitable for transfer calculations. Since the PARAMETERSET groups of Table 7.3 define values of ASYMP_TOPIA and STEPSPER that are to be used in the two-body states, the PARAMETERSET keyword should precede the definitions of the two-body states. The CA60A and CA60B PARAMETERSET names are appropriate for two-nucleon transfer reactions initiated by $^{16}$O on Ca near 60 MeV. The PB100A, PB100B, and PB100C parameter set names are designed for single-nucleon $^{16}$O on Pb reactions near the Coulomb barrier; at bombarding energies of several hundred MeV, larger values of LSTEP may be used. The ALPHA1, ALPHA2 and ALPHA3 sets were designed for ($^{16}$O, $^{12}$C) reactions on targets around $^{24}$Mg.

Table 7.3 shows that as one increases the numbers of grid points, one should also reduce DW_CUTOFF so as to include more of the integrand in the computation. This reduction in turn means that ASYMP_TOPIA may have to be made larger. The larger intervals that result from the smaller DW_CUTOFF will also result in the Gauss points being spread out further so that the same number of points will give reduced accuracy. The ALPHA1 sets have significantly larger NPSUM and NPDIF values than the other sets. These are due to the facts that the alpha-core bound state wavefunctions have a large number of nodes, and that the DIF grids for alpha transfer extend further than for than for one- and two-nucleon transfer.

It is strongly suggested that for each substantially new problem, the user make calculations with several different sets of grid parameters to verify that convergence has
been achieved. Many nodes in the bound-state wavefunctions, a strong dependence on $L$ (such as is obtained with surface transparent potentials) of the elastic scattering wavefunctions, poorly matched reactions, and transfers of heavy clusters are all examples of cases in which modifications of the parameter sets may be necessary.

7.4 Saving Time and Core with SAVEHS and USEHS

This section is retained for historical reasons. Modern computers have so much memory and also are so fast that the savings described here are irrelevant.

Almost all of the time in a large DWBA calculation is devoted to the form factor integral:

$$H(R_i, R_{out}) = \int d\phi \ (\phi_P \ V_{eff} \ \phi_T)$$

These quantities are independent of both the optical potentials and the scattering energy. Most DWBA studies are principally concerned with the effects of variations in the optical potentials. Thus it is reasonable to save the computed $H$’s in a dataset for reuse with different optical potentials. If a study is being made at a number of relatively close energies, time can also be saved by making one large calculation containing all of the orbital angular momenta needed for all of the energies, and then in subsequent jobs using subsets of the resulting dataset. PTOLEMY provides two keywords, SAVEHS and USEHS, to allow the form-factor integrals to be saved.

The SAVEHS keyword is used to initiate the saving of the integrals over $\phi$ of the form factor. These integrals may then be reused in later calculations with different optical potentials at a considerable saving in CPU time. The SAVEHS keyword must be entered before the first semi-colon and should be entered after the HEADER and REACTION keywords if they are used. The output will be written to Fortran unit 1.

The USEHS keyword is used to indicate that the $H$-integrals saved in a previous calculation with the SAVEHS keyword are to be reused. USEHS must be entered before the first semicolon. If it is used, the REACTION, LSTEP and all grid-setting keywords should not be specified again. The bound state potentials must not be entered; rather the definitions of the new optical potentials should directly follow the USEHS keyword. These definitions are then followed by the final semicolon indicating that the DWBA calculation is to begin. Thus a typical USEHS run has the form

```plaintext
USEHS
   INCOMING potential definition;
   OUTGOING potential definition;
;
RETURN
```

If ELAB, LMIN, and/or LMAX are not specified, the SAVEHS values will be used. However new values of these parameters may be specified in the USEHS run. If different values are to be specified; they must be given after the USEHS keyword. If a different LMIN is specified, the user must insure that it was one of the values explicitly computed in the SAVEHS run [i.e. that $\text{LMIN} = \text{LMIN}($SAVEHS$) + \text{n}\times\text{LSTEP}$]. If a new value of ELAB is specified, the SUM
### 7.4. SAVING TIME AND CORE WITH `SAVEHS` AND `USEHS`

<table>
<thead>
<tr>
<th>Keyword</th>
<th>Default</th>
<th>CA60A</th>
<th>CA60B</th>
<th>PB100A</th>
<th>PB100B</th>
<th>PB100C</th>
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<tr>
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<td>3</td>
<td>2</td>
<td>5</td>
<td>4</td>
<td>3</td>
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<tr>
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<td>$10^{-2}$</td>
<td>$10^{-3}$</td>
<td>$10^{-3}$</td>
<td>$3 \times 10^{-3}$</td>
<td>$10^{-4}$</td>
</tr>
<tr>
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<td>6.</td>
<td>6.</td>
<td>2.5</td>
<td>3.</td>
<td>3.5</td>
</tr>
<tr>
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<td>15</td>
<td>10</td>
<td>15</td>
<td>20</td>
</tr>
<tr>
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<td>10</td>
<td>13</td>
<td>8</td>
<td>10</td>
<td>14</td>
</tr>
<tr>
<td>NPHI</td>
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<td>10</td>
<td>12</td>
<td>10</td>
<td>12</td>
<td>16</td>
</tr>
<tr>
<td>LMINSUB</td>
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<td>10</td>
<td>15</td>
<td>12</td>
<td>16</td>
<td>20</td>
</tr>
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<td>.6</td>
<td>.5</td>
<td>.86</td>
<td>.78</td>
<td>.70</td>
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<td>30</td>
<td>35</td>
<td>40</td>
</tr>
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<td>1.6</td>
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<td>5.</td>
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<td>$10^{-3}$</td>
<td>$10^{-3}$</td>
</tr>
<tr>
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</tr>
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<td>55</td>
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<tr>
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<td>12</td>
<td>16</td>
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</table>

<table>
<thead>
<tr>
<th>Keyword</th>
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<th>ALPHA3</th>
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<tbody>
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<td>1</td>
<td>1</td>
</tr>
<tr>
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<td>$10^{-4}$</td>
<td>$10^{-5}$</td>
</tr>
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<td>7.</td>
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</tr>
<tr>
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<td>70</td>
</tr>
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<td>NPDIFF</td>
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</tr>
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<tr>
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<td>15</td>
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</tr>
<tr>
<td>LMINMULT</td>
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<td>.5</td>
<td>.4</td>
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</tr>
<tr>
<td>LMAXMULT</td>
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</tr>
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<td>3.</td>
<td>3.</td>
</tr>
<tr>
<td>GAMMADIF</td>
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</tr>
<tr>
<td>ASYMPTOPIA</td>
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<td>30</td>
</tr>
<tr>
<td>STEPSPER</td>
<td>12</td>
<td>20</td>
<td>25</td>
</tr>
</tbody>
</table>

Table 7.1: `PARAMETERSET` names and associated values for DWBA transfer calculations. The first column gives the default values.
grid will not be the optimal grid since it was chosen for a different energy. In cases in which it is desired to do USEHS runs at different energies, the SAVEHS run should be made at the maximum energy to be used. It should be made with more SUM points than would be necessary for a calculation at a single energy. Also the automatic choice of LMIN will have to be reduced.

Since the large arrays used for the angular transforms in the form-factor integral do not need to be constructed for USEHS calculations, there are also substantial memory savings in USEHS jobs. It is suggested that during a study of a given set of reactions, a SAVEHS run be made for each different reaction, and the results stored on the on-line disks. The effects of different optical potentials may then be quickly and cheaply studied for all the reactions using USEHS jobs. When the study is complete, the SAVEHS datasets may be copied to a tape for long-term storage.

### 7.5 Reading the Output

As each of the bound states and elastic scattering states are entered, a summary of the two-body channel is printed; the format of this summary has been described in Sec. 4.5. When the fifth semicolon is entered, a summary of the reaction is printed. This summary lists the nuclei involved, the bound-state properties, and the spectroscopic factors. The \( Q \)-value of the reaction is also printed. If the \( Q \)-value or the bound-state or outgoing scattering energies were explicitly entered by the user, then the differences of the bound-state energies and of the scattering energies may not be the same, or they may not be equal to the \( Q \)-value. Under these circumstances an appropriate warning message is printed, however the calculation proceeds with the bound-state and scattering energies that were listed in the previous summaries. After the bound-state properties, the content of the effective interaction, as determined by one of the keywords NUONLY, USESIMPCOUL, USECOULOMB, or USECORE, is listed.

The range of \( L \)'s and the increment of \( L \) for which the radial integrals will be explicitly computed is then listed. The range of transferred orbital angular momenta is also given.

A several-line summary of the three-dimensional integration grid is then given. The lines give the number of Gauss points used for, and the extent of, each dimension of the grid. The entry under NUM. PTS for \((RI + RO)/2\) gives the number of Gauss points used in the integrals involving the wavefunctions, and is determined by the value of SUMPOINTS; the form factors are found at these points by interpolation. The number of points in the \((RI + RO)/2\) grid at which the form factors are evaluated (NPSUM) is given in parentheses at the end of the \((RI + RO)/2\) line. The warning message

\[
\text{INVALID VMIN, VMID, VMAX...}
\]

may precede the integration grid summary; it is explained in Sec. 7.2. Two lines of the form

\[
\text{MAXIMUM R'S USED FOR...}
\]

give the maximum radial coordinate values for which each of the bound-state and scattering-state wavefunctions will be used in the three-dimensional integrals. The corresponding values of ASYMPTOPIA may be reduced to approximately 5 fm greater than these values
7.5. **READING THE OUTPUT**

in future calculations. If the bound state **ASYMPTOPIA** prove to be limiting factors in the grid construction, the warning message

**ERROR ... BOUND STATE WAVEFUNCTIONS NEEDED...**

 will be printed. The calculation should be repeated with an increased value of **ASYMPTOPIA** for the bound states since the radial integrals computed with the smaller grid may not be accurate. The increased value should be at least 5 fm larger than the values printed in the warning message.

If (as is suggested) PRINT=2 has been specified, one or more pages labeled **COMPUTATION OF RADIAL INTEGRALS** will next be printed. These pages contain only those radial integrals that are explicitly computed; the radial integrals for even Lout are printed first. Three columns give the real and imaginary parts and the magnitude of the radial integrals. These are followed by two columns labeled **CANCELLATIONS** the second of which is not presently used. The first column gives an indication of the numerical cancellations that occurred in the \( dR_i dR_o \) integral and is defined as

\[
\frac{\int dR_i dR_o R_i R_o |f(out)Hf(in)|}{\int dR_i dR_o R_i R_o |f(out)Hf(in)|} \tag{7.2}
\]

where the denominator is just the magnitude of the radial integral. The entries in this column are usually quite large (>100) for the smaller values of \( L \), since these radial integrals are small by virtue of large cancellations. However, the values for \( L \geq L_{crit} \) are usually less than 5, indicating that there was not much difficulty associated with these radial integrals. If all of the entries in this column are greater than 10, then the reaction is in some way poorly matched, and the convergence of the computed cross sections should be carefully checked.

The page labeled **INTERPOLATION AND EXTRAPOLATION IN L** gives a summary of the extrapolation of the radial integrals from LMAX to \( L'_{\text{max}} \). The value of \( L'_{\text{max}} \) is printed in the line **MAXIMUM LO USED...**. This line may be preceded or followed by a number of warning or error messages indicating difficulties in the extrapolation to \( L'_{\text{max}} \). These difficulties are usually associated with the fact that some \( (L_x, L_{in}-L_{out}) \) combinations are poorly matched and the computed radial integrals contain large errors. If the errors are large enough, the computed radial integrals will not have the correct asymptotic form, and it will be impossible to fit the extrapolating function to them. If the extrapolated radial integrals have a negligible effect on the computed differential cross sections, then these error messages may be ignored. An indication of this may be found in the table labeled **SUMMARY OF EXTRAPOLATION PARAMETERS** that is printed at the end of this page. The \( (L_x, L_{in}-L_{out}) \) combinations that could not be extrapolated have zeros in the columns that give the extrapolation parameters. The columns labeled **PEAK** give the Lout for which the radial integrals with the printed values of \( L_x \) and \( L_{in}-L_{out} \) have their maximum magnitude and that maximum magnitude. If the maximum magnitude is small compared to other maximum magnitudes, then the extrapolated values (which are usually at least a factor 10 smaller) are probably negligible and the failure to be able to extrapolate may be safely ignored.

The pages labeled **REACTION AND ELASTIC PARTIAL...** give all of the radial integrals
and the elastic $S$-matrix elements for $L_{\text{MIN}} \leq L \leq L_{\text{MAX}}$. The format of these pages was explained in Sec 6.3.

Finally the cross sections are given on the pages labeled \textsc{Computations of Cross Sections}. Most of the entries on these pages were also explained in Sec. 6.3, and only the differences will be given here. The column labeled \% FROM $L > L_{\text{MAX}}$ gives (as a percentage) the contribution to the cross section from the radial integrals for $L_{\text{MAX}} < L \leq L'_{\text{MAX}}$. Our experience is that this contribution is usually accurate to somewhat better than 10\%. Thus if at a given angle the entry is 5\%, one may assume that the extrapolation procedure introduced an error of less than 0.5\% to the differential cross section. If more than one $L_x$ can contribute to the reaction, there will be columns giving the differential cross section for each $L_x$; the cross section in the second column is the sum of these partial cross sections. Of course the spectroscopic factors are included in all cross sections.
Chapter 8

Control Keywords

In this section we describe some of the control keywords that are available in PTOLEMY. These keywords may generally be placed anywhere in the PTOLEMY input and they usually cause some sort of immediate action; they are not followed by a semi-colon. These keywords are listed more or less in decreasing order of interest; all of them except RETURN are optional.

The RETURN or END keyword should be used as the last keyword in every PTOLEMY job. It causes allocator statistics to be printed and then returns control to the operating system.

The HEADER keyword may be used to enter a header that will be printed on the top of most pages of the PTOLEMY output. The header will consist of the remainder of the input line beginning with the first alphanumeric following the HEADER keyword. In addition the REACTION specification (if it is entered) and the laboratory energy will always be part of the header.

The KEEP keyword is used to keep PTOLEMY results in a form suitable for later recovery with the SPEAKEASY KEPT command. KEEP must be followed by two names: the first is the PTOLEMY name of the item to be kept and the second is the name under which it is to be saved in the current directory (with an operating-system dependent suffix). This second name must be different from all other names already in the directory or else the new object will replace the previously kept object. The following is a list of the PTOLEMY names of the objects the user is most likely to want to keep:

- **ANGLEGRID** - A 3-element array containing ANGLESTEP, ANGLEMIN and ANGLEMAX.

- **CROSSSEC** - The differential cross section on the (ANGLEMIN, ANGLESTEP, ANGLEMAX) grid. For DWBA calculations this contains the reaction cross section. For stand-alone elastic scattering it contains the elastic differential cross sections. In both cases it is in mb/sr.

- **TORUTHERFORD** - The ratio of the elastic scattering differential cross section to the Rutherford cross section. This array is produced only in stand-alone elastic scattering.
- **LXCROSSSECTION** - A Num(angles) × Num($L_x$) array containing the DWBA cross sections for each $L_x$ (summed over $M_x$) at each angle.

- **MXCROSSSECTION** - A Num(angles) × Num($L_i$, $L_x$) array containing the DWBA cross sections for each ($L_x$, $M_x$) at each angle. The order of the columns is

$$
(L_x, M_x) = (L_{x,\text{min}}, 0), (L_{x,\text{min}}, 1), \ldots, (L_{x,\text{min}}, L_{x,\text{min}}),
(L_{x,\text{min}} + 1, 0), \ldots, (L_{x,\text{max}}, L_{x,\text{max}})
$$

(8.1)

This array is available to **KEEP** only if the keyword `SAVEMXCROSS` (see below) has been entered.

- **SMAG** and **SPHASE** - The magnitude and phase of the S-matrices. These two objects are each Num($L$) × Num($L_i$, $L_x$) arrays. The arrangement of the columns is such that $S(L_x, L_i, L_{out})$ is indexed as SMAG(j,k) and SPHASE(j,k) with

$$
j = L_{out} - L_{\text{MIN}} + 1
$$

$$
k = [(L_x + L_{x,\text{min}} + 1)(L_x - L_{x,\text{min}}) + L_x + L_i - L_{out}] / 2 + 1
$$

- **B** - A Num(angles) × Num($L_i$, $L_x$) complex array containing the $B(L_x, M_x, \theta)$ of Eq ??? or 2.55 for each ($L_x$, $M_x$) at each angle. The columns have the same order as for **MXCROSSSECTION**. This array is available for **KEEP** only if the `SAVEB` keyword (see below) has been entered.

- **SIN** - The elastic S-matrix elements in the incoming channel.

- **SOUT** - The elastic S-matrix elements in the outgoing channel.

- **WAVER, WAVEI** - The real and imaginary parts of the most recently computed elastic scattering wavefunction.

- **PHIn** - The n’th bound state wavefunction.

- **HEADER** - A character array (up to 65 characters) containing the **HEADER**.

- **REACTION** - A character array (up to 45 characters) containing the **REACTION** or **CHANNEL** specification.

- **ELAB** - A scalar with the laboratory energy.

The following symbols were used in the above definitions of the objects for the **KEEP** command:

- **$L_{x,\text{max}}$** - Maximum $L_x$ (transfered $L$)

- **$L_{x,\text{min}}$** - Minimum $L_x$

- **Num(angles)** - Number of angles
- Num($L$) - $L_{\text{max}} - \text{LMIN} + 1$

- Num($L_x$) - $L_{x,\text{max}} - L_{x,\text{min}} + 1$

- Num($L_i, L_x$) - $[(L_{x,\text{max}} + L_{x,\text{min}} + 2)(L_{x,\text{max}} - L_{x,\text{min}} + 1)]/2$ (This expression does not take into account abnormal parity restrictions but it is always the correct expression to use.)

The SAVEMXCROSS keyword may be entered to cause the DWBA cross sections for every ($L_x$, $M_x$) and angle to be saved. The SAVEB keyword may be entered to cause the $B(L_x, M_x, \theta)$ to be saved for every ($L_x$, $M_x$) and angle. The resulting objects (MXCROSSSECTION and B) may then be stored using the KEEP command. The defaults are NOSAVEMXCROSS and NOSAVEB.

The WRITENS command is an alternative method of storing objects that are in the PTOLEMY allocator for use by other programs. The command is followed by the name or number of an object in the allocator. The object will be written on Fortran unit 15 using a (1X, 4G17.8) format. The names and structures of the available objects are given in the description of the KEEP command above, however two-dimensional objects will be written as the transpose of the Speakeasy formats given above.

The NEWPAGE keyword may be used to cause subsequent PTOLEMY output to begin on a new page. It will probably be used only if several stand-alone two-body calculations are being done.

The KEYWORDS keyword may be used at any time to cause a listing of the current settings of all data-entering and option-choosing keywords. It is suggested that it be used at the end of all runs to provide a verification of the parameters and options in effect. The LISTKEYS keyword will list the names of all valid keywords.

The UNDEFINE keyword may be used to set the status of a keyword to “undefined.” It is followed by the name of the keyword that is to be undefined. The most likely use of UNDEFINE is to undefine a potential radius ($R$, $R_I$, etc.) during stand-alone calculations so that it will automatically be computed from the corresponding radius parameter ($R_0$, $R_I0$, etc.) the next time it is needed. One may also want to undefine STEPSPER after the use of PARAMETERSET so that a specific STEPSIZE may be entered.

The NSCATALOG keyword may be used at any time to provide a list of the names and sizes of all currently defined objects in the allocator. It will also give the NSSTATUS output.

The NSSTATUS keyword will cause a short summary of the allocator status to be printed. This summary will show the allocator size, its current in-use size and the peak in-use size. This summary is automatically printed at the end of all PTOLEMY jobs.

The SIZE keyword may be used to specify in some versions of PTOLEMY the size of the allocator (the PTOLEMY work area); it is the size in bytes of the allocator. If it is to be used, the SIZE keyword should be on the first line of PTOLEMY input.

The DUMP or NSDUMP keywords may be used to print an object in the allocator. They are followed by the name or number of the object to be printed.

The RESET keyword may be used to cause PTOLEMY to be set back to its initial status. The complete allocator is cleared and all keywords are set to their default values.
However the allocator size may not be changed after a \texttt{RESET}. The use of \texttt{RESET} allows several independent calculations to be made in one job. If it is desired to do a \texttt{SAVEHS} calculation and immediately follow it with several \texttt{USEHS} calculations, each \texttt{USEHS} step should be preceded with a \texttt{RESET}.

The \texttt{CLEAR} keyword causes the allocator to be cleared. All objects are removed from the allocator. However none of the keyword settings are changed. The size of the allocator may not be respecified after a \texttt{CLEAR} command.

The \texttt{COPY} keyword may be used to copy the data in an object in the allocator into a second object. The form of the command is

\begin{verbatim}
COPY fromname toname
\end{verbatim}

where “fromname” must be the name of an object already in the allocator. If “toname” already exists in the allocator and is the same size as “fromname,” the data in “fromname” will be copied into “toname.” Otherwise “toname” will be created (or changed to have the correct size) and then the copy will occur. In all cases “fromname” is not changed.

The \texttt{COPY} command may be used to “fool” \textsc{ptolemy} into using a different potential or wavefunction in a subsequent part of the calculation.

The \texttt{RENAME} command is used to change the name of an object in the allocator. Its form is

\begin{verbatim}
RENAME oldname newname
\end{verbatim}

where “oldname” must be the name of an object in the allocator. The name of this object will be changed to “newname.” There must not be another object with the name “newname” already in the allocator; if there is, inconsistent results may occur.

The \texttt{FREE} command is used to delete an object from the allocator and thus make its space available for other objects. The command is followed by the name of the object to be freed. If the object does not exist, a warning will be printed and processing will continue with the next input line.

In the \texttt{WRITENS}, \texttt{COPY}, \texttt{RENAME} and \texttt{FREE} commands the first object name may be replaced with the actual number of the object in the allocator. However this practice is not recommended since it is difficult to predict the numbers of the objects.

An object may be added to or changed in the allocator by means of the \texttt{ALLOCATE} command. This command has the form

\begin{verbatim}
ALLOCATE name list-of-numbers
\end{verbatim}

where “name” will be the name of the object in the allocator. The number of elements in the list-of-numbers determines the length of the object. The numbers may be in any format, with as many or as few as is desired per input line. If an object with the name “name” already exists, it is replaced with the new object as defined by the \texttt{ALLOCATE} command. In this manner one may read in arbitrary bound-state wavefunctions after \textsc{ptolemy} has computed the bound-state wavefunctions for a Woods-Saxon potential. The new wavefunctions will then be used in a subsequent DWBA calculation.

The \texttt{BIMULT} keyword causes two objects (they may be the same object) to be multiplied together. The form of the command is

\begin{verbatim}
BIMULT name1 name2
\end{verbatim}

where “name1” and “name2” are the names of the two arrays to be multiplied together. They must be of the same length. They will be multiplied together in an element-by-
element fashion and the resulting array will be stored in a new unnamed object in the allocator. Its number will be printed in the output. The numbers of the input arrays may be used inplace of their names.

The keyword \texttt{NUMRNUM} may be used to compute the matrix element between two wavefunctions of a power of $r$. The form of the command is

\begin{verbatim}
NUMRNUM name1 power name2
\end{verbatim}

where “name1” and “name2” are the names (or numbers) of two objects in the allocator. They must be of the same length and must be wavefunctions that were computed with the present value of \texttt{STEPSIZE}. The integer “power” is the power of $r$ that is to be included in the integral. The integral

\begin{equation}
\int_{\text{ASYMPTOPIA}}^{\infty} drr^{\text{power}} \times \text{name1} \times \text{name2}
\end{equation}

will be computed and printed.

The keyword \texttt{NRNLIMS} may be used to compute the partial matrix element of a power of $r$. The form of the command is

\begin{verbatim}
NRNLIMS name1 power name2 start stop
\end{verbatim}

where “name1”, “name2,” and “power” are the same as for \texttt{NUMRNUM}. The range of the integral is specified by “start” and “stop” which are specified in fm.

The keyword \texttt{PRINT} may be used to control the amount of printing that PTOLEMY does. It is followed by a five-digit integer that indicates the amount of printing that is to occur. Each digit controls different items of the printed output. The larger the digit, the more information that is printed. The default value is $\text{PRINT} = 10001$ which results in summaries of the input, the radial integral phases and magnitudes and the cross sections being printed. If the five-digit number is written as $\text{PRINT} = \text{TMCXI}$, the significance of the digits is:
CHAPTER 8. CONTROL KEYWORDS

I = 0 - Only print the differential cross sections and final fit values.
1 - (Default) - Print summaries of input, magnitudes and phases of the radial integrals and elastic S-matrices in addition to output for PRINT=0. For fits the initial and final values are shown along with a summary of the path followed by the fitter.
2 - Print radial integrals as they are computed and give estimates of their cancellations. This option is strongly recommended as large cancellations are an indication that the convergence should be checked. The singular values are printed by the QUAVER fitter.

≥ 3 - Print debugging information.

X = 1 - The WKB amplitudes used to find the critical L are printed for each L.
2 - Debugging output is printed by the WKB routine.

C = 1 - Several lines are printed for every chi-squared function calculation made during the course of a fit. Debugging output from the Coulomb excitation integrals is printed. Work arrays are not freed following the radial integral computation.
2 - Convergence to the bound-state eigenvalue is printed.
4 - Debugging output from the L-interpolation is printed.

M = 1 - The elastic S-matrix element is printed every time a scattering wavefunction is computed.
4 - Debugging output is produced by the elastic wavefunction routine.

T = 1 - (Default) - Show conversion of FIT data to standard form.
4 - Some debugging output about the transfer effective interaction is produced.
9 - Enormous amounts of debugging output from the transfer effective interaction (the end has never been seen) are produced.

The value of PRINT may be changed at any time to effect subsequent printing except that in DWBA calculations the value of PRINT that was in effect at the time of the specification of the outgoing scattering state will determine the printing of the elastic S-matrices during the computation of the radial integrals.
Chapter 9

Acknowledgments

Much of the original development of PTOLEMY was done by David Gloeckner. We wish to thank Frank Serduke for making available the subroutines that supply the data on nuclear charges, mass excesses, and spins. Larry Nazareth has been most helpful in giving us advance access to a number of subroutines that are being considered for inclusion in the Argonne Applied Mathematics Division MINPACK project. The Coulomb function subroutine is largely based on the Manchester subroutine RCWFN.¹

Appendix A

List of all Keywords

The following lists contain brief descriptions and default values for all the PTOLEMY keywords. Some of these keywords have not been discussed in the above text either because they are primarily designed for use in debugging PTOLEMY or because it is doubtful that the average user will need them. They are all included here for completeness. The entry “none” under “Default” means that the keyword is initially undefined. If it is not defined in the input and is necessary to the calculation, an error message will be printed and the job aborted. The entry “none*” under “Default” means that the keyword is initially undefined, but if it is not defined in the input, an appropriate value will be found by PTOLEMY.

The keywords CROSSSECTION, GRIDSETUP, LINTERPOL, and RADIALINT are used in place of the final semicolon to initiate individual stages of the DWBA calculation and should not be used in standard calculations. If desired the keywords DWBA or NZRDWBA may be used before the final semicolon to indicate that a DWBA calculation is to be done but they are not, at present, necessary.

All lengths are specified in fm and all energies are in MeV.
### A.1 Reaction-defining Keywords

<table>
<thead>
<tr>
<th>Keyword</th>
<th>Default</th>
<th>Meaning</th>
</tr>
</thead>
<tbody>
<tr>
<td>BETA</td>
<td>none</td>
<td>Nuclear deformation parameter</td>
</tr>
<tr>
<td>BETACOULOMB</td>
<td>none</td>
<td>Coulomb deformation parameter</td>
</tr>
<tr>
<td>BELX</td>
<td>none</td>
<td>$B (E, L_x, \text{excitation})$</td>
</tr>
<tr>
<td>E*i</td>
<td>0</td>
<td>Excitation energy in MeV of nucleus $i$ ($i = A, B, \text{BIGA}, \text{BIGB}, X$)</td>
</tr>
<tr>
<td>ECM</td>
<td>none</td>
<td>Incoming c.m. scattering energy in MeV</td>
</tr>
<tr>
<td>ELAB</td>
<td>none</td>
<td>Incoming Laboratory scatter energy in MeV</td>
</tr>
<tr>
<td>Ji</td>
<td>none</td>
<td>Intrinsic spin of nucleus $i$</td>
</tr>
<tr>
<td>Mi</td>
<td>none</td>
<td>Mass in AMU of nucleus $i$ ($i = A, B, \text{BIGA}, \text{BIGB}, X$)</td>
</tr>
<tr>
<td>MXCi</td>
<td>none</td>
<td>Total mass excess in MeV of nucleus $i$ ($i = A, B, \text{BIGA}, \text{BIGB}, X$)</td>
</tr>
<tr>
<td>MXCGi</td>
<td>none</td>
<td>Ground state mass excess in MeV of nucleus $i$ ($i = A, B, \text{BIGA}, \text{BIGB}, X$)</td>
</tr>
<tr>
<td>NUONLY</td>
<td>off</td>
<td>Use only the nuclear part of the B.S. potential in the interaction potential</td>
</tr>
<tr>
<td>Q</td>
<td>none</td>
<td>$Q$-value in MeV</td>
</tr>
<tr>
<td>REACTION</td>
<td>none</td>
<td>Defines the reaction in standard notation</td>
</tr>
<tr>
<td>SPAMP</td>
<td>1</td>
<td>Projectile spectroscopic amplitude</td>
</tr>
<tr>
<td>SPAMT</td>
<td>1</td>
<td>Target spectroscopic amplitude</td>
</tr>
<tr>
<td>SPFACP</td>
<td>1</td>
<td>Projectile spectroscopic factor</td>
</tr>
<tr>
<td>SPFACT</td>
<td>1</td>
<td>Target spectroscopic factor</td>
</tr>
<tr>
<td>USECORE</td>
<td>on</td>
<td>Use Coulomb and real nuclear core corrections in the interaction potential</td>
</tr>
<tr>
<td>USECOULOMB</td>
<td>off</td>
<td>Use Coulomb with core corrections in the interaction potential</td>
</tr>
<tr>
<td>USEPROJECTILE</td>
<td>on</td>
<td>Use projectile B.S. potential as the interactional potential</td>
</tr>
<tr>
<td>USESIMPCOULOMB</td>
<td>off</td>
<td>Use Coulomb of just one B.S. in the interaction potential</td>
</tr>
<tr>
<td>USETARGET</td>
<td>off</td>
<td>Use target B.S. potential as the interaction potential</td>
</tr>
<tr>
<td>Zi</td>
<td>none</td>
<td>Charge of nucleus $i$ ($i = A, B, \text{BIGA}, \text{BIGB}, X$)</td>
</tr>
</tbody>
</table>
### A.2 Two-body State Keywords

<table>
<thead>
<tr>
<th>Keyword</th>
<th>Default</th>
<th>Meaning</th>
</tr>
</thead>
<tbody>
<tr>
<td>A</td>
<td>none</td>
<td>Real W.S. diffuseness</td>
</tr>
<tr>
<td>AE</td>
<td>0</td>
<td>$AE*ELAB$ is added to $A$</td>
</tr>
<tr>
<td>AESQ</td>
<td>0</td>
<td>$AESQ*ELAB^2$ is added to $A$</td>
</tr>
<tr>
<td>AI</td>
<td>none</td>
<td>Imaginary W.S. diffuseness</td>
</tr>
<tr>
<td>AIE</td>
<td>0</td>
<td>$AIE*ELAB$ is added to $AI$</td>
</tr>
<tr>
<td>AIESQ</td>
<td>0</td>
<td>$AIESQ*ELAB^2$ is added to $AI$</td>
</tr>
<tr>
<td>ASI</td>
<td>none</td>
<td>Surface absorption diffuseness</td>
</tr>
<tr>
<td>ASO</td>
<td>none</td>
<td>Real spin-orbit diffuseness</td>
</tr>
<tr>
<td>ASOI</td>
<td>none</td>
<td>Imaginary spin orbit diffuseness</td>
</tr>
<tr>
<td>CHANNEL</td>
<td>none</td>
<td>Specifies a two-body channel</td>
</tr>
</tbody>
</table>
| E       | none    | Two-body c. m. energy in MeV  
|         |         | ($ELAB$ or $ECM$ may be used in stand-alone or in fit) |
| EINVERSE| off     | Energy-dependent pots are in $1/ELAB$ |
| EPOWERS | on      | Energy-dependent pots are in $ELAB$ |
| E*P     | none    | Projectile excitation energy |
| E*T     | none    | Target excitation energy |
| J       | none    | Total two-body angular momentum |
| JP      | none    | Total projectile angular momentum (for spin-orbit) |
| I       | none    | Orbital angular momentum |
| M       | none    | Reduced mass in Mev/c^2 |
| MP      | none    | Projectile mass in AMU |
| MT      | none    | Target mass in AMU |
| NODES   | none    | Number of bound state nodes for $r > 0$. |
| R       | none    | Real W.S. radius. |
| R0      | none    | Real W.S. radius parameter |
| ROE     | 0       | $ROE*ELAB$ is added to $R0$ |
| ROESQ   | 0       | $ROESQ*ELAB^2$ is added to $R0$ |
| RC      | none    | Coulomb radius |
| RCO     | none    | Coulomb radius parameter |
| RCP     | none    | Coulomb radius of projectile |
| RCT     | none    | Coulomb radius of target |
| RCOP    | none    | Coulomb radius parameter of projectile |
| RCOT    | none    | Coulomb radius parameter of target |
| RI      | none    | Imaginary W.S. radius |
| RIO     | none    | Imaginary W.S. radius parameter |
| RIOE    | 0       | $RIOE*ELAB$ is added to $RIO$ |
| RIOESQ  | 0       | $RIOESQ*ELAB^2$ is added to $RIO$ |
| RSI     | none    | Surface absorption radius |
| RSIO    | none    | Surface absorption radius parameter |
| RSO     | none    | Real spin-orbit radius |
| RSOO    | none    | Real spin-orbit radius parameter |
### A.2. TWO-BODY STATE KEYWORDS

<table>
<thead>
<tr>
<th>Keyword</th>
<th>Default</th>
<th>Meaning</th>
</tr>
</thead>
<tbody>
<tr>
<td>RSOI</td>
<td>none</td>
<td>Imaginary spin-orbit radius</td>
</tr>
<tr>
<td>RSOI0</td>
<td>none</td>
<td>Imaginary spin-orbit radius parameter</td>
</tr>
<tr>
<td>SP</td>
<td>none</td>
<td>Projectile spin</td>
</tr>
<tr>
<td>SPAM</td>
<td>none</td>
<td>Channel spectroscopic amplitude</td>
</tr>
<tr>
<td>ST</td>
<td>none</td>
<td>Target spin</td>
</tr>
<tr>
<td>TAU</td>
<td>0</td>
<td>Real S.O. depth relative to real W.S. depth</td>
</tr>
<tr>
<td>TAU1</td>
<td>0</td>
<td>Imaginary S.O. depth relative to imaginary W.S. depth</td>
</tr>
<tr>
<td>V</td>
<td>0</td>
<td>Real W.S. well depth</td>
</tr>
<tr>
<td>VE</td>
<td>0</td>
<td>$VE \times \text{ELAB}$ is added to $V$</td>
</tr>
<tr>
<td>VESQ</td>
<td>0</td>
<td>$VESQ \times \text{ELAB}^2$ is added to $V$</td>
</tr>
<tr>
<td>VI</td>
<td>0</td>
<td>Imaginary W.S. well depth</td>
</tr>
<tr>
<td>VIE</td>
<td>0</td>
<td>$VIE \times \text{ELAB}$ is added to $VI$</td>
</tr>
<tr>
<td>VIESQ</td>
<td>0</td>
<td>$VIESQ \times \text{ELAB}^2$ is added to $VI$</td>
</tr>
<tr>
<td>VSI</td>
<td>0</td>
<td>Surface absorption strength</td>
</tr>
<tr>
<td>VSO</td>
<td>0</td>
<td>Real S.O. well depth</td>
</tr>
<tr>
<td>VSOI</td>
<td>0</td>
<td>Imaginary S.O. well depth</td>
</tr>
<tr>
<td>ZP</td>
<td>none</td>
<td>Projectile charge</td>
</tr>
<tr>
<td>ZT</td>
<td>none</td>
<td>Target charge</td>
</tr>
</tbody>
</table>
### A.3 Calculation Keywords

<table>
<thead>
<tr>
<th>Keyword</th>
<th>Default</th>
<th>Meaning</th>
</tr>
</thead>
<tbody>
<tr>
<td>ACCURACY</td>
<td>$10^{-2}$</td>
<td>Accuracy of bound state convergence</td>
</tr>
<tr>
<td>ANGLEMAX</td>
<td>90</td>
<td>Maximum scattering angle in degrees</td>
</tr>
<tr>
<td>ANGLEMIN</td>
<td>0</td>
<td>Minimum scattering angle in degrees</td>
</tr>
<tr>
<td>ANGLESTEP</td>
<td>1</td>
<td>Scattering angle increment in degrees</td>
</tr>
<tr>
<td>ASYMPHIEA</td>
<td>20</td>
<td>Start of asymptotic region</td>
</tr>
<tr>
<td>COULLBMBIFT</td>
<td>1</td>
<td>Determines start of Belling expansion</td>
</tr>
<tr>
<td>DATA</td>
<td>–</td>
<td>Enters experimental data for a fit</td>
</tr>
<tr>
<td>DELTAVK</td>
<td>0.05</td>
<td>Bound state search step size</td>
</tr>
<tr>
<td>DERIVSTEP</td>
<td>$10^{-6}$</td>
<td>Stepsize for numeric gradient of potentials</td>
</tr>
<tr>
<td>DWCUTOFF</td>
<td>$10^{-3}$</td>
<td>Minimum relative integrand to use in DWBA integral;</td>
</tr>
<tr>
<td></td>
<td></td>
<td>minimum relative Coulomb amplitude for excitation</td>
</tr>
<tr>
<td>FITACCURACY</td>
<td>$10^{-3}$</td>
<td>Required relative accuracy of optical parameter fit</td>
</tr>
<tr>
<td>FITMULTIPLE</td>
<td>500</td>
<td>Meaning depends on optical model fitter in use</td>
</tr>
<tr>
<td>FITRATIO</td>
<td>$10^{-4}$</td>
<td>Meaning depends on optical model fitter in use</td>
</tr>
<tr>
<td>GAMMADIF</td>
<td>5</td>
<td>DIF compression parameter</td>
</tr>
<tr>
<td>GAMMASUM</td>
<td>1</td>
<td>SUM compression parameter</td>
</tr>
<tr>
<td>INELASACC</td>
<td>$10^{-5}$</td>
<td>Required accuracy of Coulomb excitation</td>
</tr>
<tr>
<td>LBACK</td>
<td>none*</td>
<td>LMAX-LBACK is start of $L$-extrapolation</td>
</tr>
<tr>
<td>LMAX</td>
<td>none*</td>
<td>Maximum scattering of partial wave $L$</td>
</tr>
<tr>
<td>LMAXADD</td>
<td>30</td>
<td>$L_{MAX} = \text{max}(L_{MAXADD} + L_{C})$</td>
</tr>
<tr>
<td>LMAXMULT</td>
<td>1.6</td>
<td>$L_{MAXMULT} \times L_{C}$</td>
</tr>
<tr>
<td>LMIN</td>
<td>none*</td>
<td>Minimum scattering partial wave $L$</td>
</tr>
<tr>
<td>LMINMULT</td>
<td>0.6</td>
<td>$L_{MIN} = \text{min}(L_{MINMULT} + L_{C})$</td>
</tr>
<tr>
<td>LMINSUB</td>
<td>20</td>
<td>$L_{C} - L_{MINSUB}$</td>
</tr>
<tr>
<td>LOOKSTEP</td>
<td>250</td>
<td>Number of steps in grid-searching for $\phi$</td>
</tr>
<tr>
<td>LSTEP</td>
<td>1</td>
<td>Increment of $L$ in radial integral computations.</td>
</tr>
<tr>
<td></td>
<td></td>
<td>In between values are found by interpolation.</td>
</tr>
<tr>
<td>LX</td>
<td>none</td>
<td>Exchanged orbital angular momentum</td>
</tr>
<tr>
<td>MAPDIFF</td>
<td>1</td>
<td>Gauss-point map type for DIF grid</td>
</tr>
<tr>
<td>MAPSUM</td>
<td>2</td>
<td>Gauss-point map type for SUM grid</td>
</tr>
<tr>
<td>MAXCOULITER</td>
<td>80</td>
<td>Max iterations for Belling’s expansion</td>
</tr>
<tr>
<td>MAXFUNCTIONS</td>
<td>50</td>
<td>Maximum number of chi. sgr. computations allowed in an optical fit mode</td>
</tr>
<tr>
<td>MAXITER</td>
<td>10</td>
<td>Max. number of iterations in bound state search</td>
</tr>
<tr>
<td>MAXLEXTRAP</td>
<td>100</td>
<td>Maximum allowed $L$-extrapolation</td>
</tr>
<tr>
<td>MIDMULT</td>
<td>2</td>
<td>SUMMID multiplier</td>
</tr>
<tr>
<td>NAIITKEN</td>
<td>4</td>
<td>degree of interpolation polynomial</td>
</tr>
<tr>
<td>NBACK</td>
<td>4</td>
<td>Num. of backward steps in scattering wave asymptotic matching</td>
</tr>
<tr>
<td>NCOSINE</td>
<td>256</td>
<td>Size of fast-cosine table</td>
</tr>
<tr>
<td>NPCOULOMB</td>
<td>8</td>
<td>Num. points/cycle in Coulomb excitation</td>
</tr>
</tbody>
</table>
### A.3. Calculation Keywords

<table>
<thead>
<tr>
<th>Keyword</th>
<th>Default</th>
<th>Meaning</th>
</tr>
</thead>
<tbody>
<tr>
<td>NPDIFF</td>
<td>10</td>
<td>Num. of points in DIF grid</td>
</tr>
<tr>
<td>NPHIADD</td>
<td>4</td>
<td>Num. of steps to extend ( \phi ) grid</td>
</tr>
<tr>
<td>NPDIFF</td>
<td>10</td>
<td>Num. of points in ( \phi ) grid</td>
</tr>
<tr>
<td>NPSUM</td>
<td>15</td>
<td>Num. of computed points in SUM grid</td>
</tr>
<tr>
<td>NUMRANDOM</td>
<td>0</td>
<td>Num. of random steps for some fitters</td>
</tr>
<tr>
<td>REINITIALIZ</td>
<td>0</td>
<td>Number of iterations between reinitialization of the fitter (mainly for POWELL65)</td>
</tr>
<tr>
<td>STEPSIZE</td>
<td>0.1</td>
<td>Step size for solutions of two-body differential equations</td>
</tr>
<tr>
<td>STEPSPER</td>
<td>none</td>
<td>Number of steps per “wavelength” for solutions of two body equations</td>
</tr>
<tr>
<td>Step1R</td>
<td>1</td>
<td>Starting value for scattering state differential equations</td>
</tr>
<tr>
<td>Step1I</td>
<td>1</td>
<td>Starting value for scattering state differential equations</td>
</tr>
<tr>
<td>SUMMAX</td>
<td>none*</td>
<td>End of SUM grid</td>
</tr>
<tr>
<td>SUMMIN</td>
<td>none*</td>
<td>Beginning of SUM grid</td>
</tr>
<tr>
<td>SUMPOINTS</td>
<td>6</td>
<td>Num. points/wavelength for SUM grid</td>
</tr>
<tr>
<td>VPOLYORDER</td>
<td>3</td>
<td>Order of interpolating poly for DIF grids</td>
</tr>
</tbody>
</table>
## Appendix A. List of All Keywords

### A.4 Control and Option-selecting Keywords

<table>
<thead>
<tr>
<th>Keyword</th>
<th>Default</th>
<th>Meaning</th>
</tr>
</thead>
<tbody>
<tr>
<td>ALLOCATE</td>
<td>-</td>
<td>Enter data into the allocator</td>
</tr>
<tr>
<td>BATCH</td>
<td>none*</td>
<td>Enter batch mode – an error terminates things</td>
</tr>
<tr>
<td>BIMULT</td>
<td>-</td>
<td>Multiply two arrays together</td>
</tr>
<tr>
<td>BOUNDSTATE</td>
<td>off</td>
<td>Stand-alone bound state is being entered</td>
</tr>
<tr>
<td>CHECKASYMP</td>
<td>off</td>
<td>Check approach of scattering state to asymptotic form</td>
</tr>
<tr>
<td>CLEAR</td>
<td>-</td>
<td>Clear allocator of all defined objects</td>
</tr>
<tr>
<td>CMANGLES</td>
<td>on</td>
<td>Angle grid is in c.m.</td>
</tr>
<tr>
<td>COMPLEXWAVE</td>
<td>on</td>
<td>Scattering waves have complex phase</td>
</tr>
<tr>
<td>COPY</td>
<td>-</td>
<td>Copy one object in the allocator into another</td>
</tr>
<tr>
<td>CROSSSECTION</td>
<td>off</td>
<td>Entering data for DWBA cross section stage</td>
</tr>
<tr>
<td>DAVIDON</td>
<td>off</td>
<td>Use the Davidon variable-metric fitter</td>
</tr>
<tr>
<td>DERIVCHECK</td>
<td>off</td>
<td>Check the analytic chi. sgr. gradients</td>
</tr>
<tr>
<td>DOASYMPT</td>
<td>on</td>
<td>Use Whittaker bound state asymptotic form</td>
</tr>
<tr>
<td>CALLTSO</td>
<td>–</td>
<td>Enter the TSO Command mode</td>
</tr>
<tr>
<td>DUMP</td>
<td>–</td>
<td>Print an object in the allocator</td>
</tr>
<tr>
<td>DUMPALL</td>
<td>–</td>
<td>The entire allocator is printed</td>
</tr>
<tr>
<td>DUMP2</td>
<td>–</td>
<td>Print an INTEGER (2) object in the allocator</td>
</tr>
<tr>
<td>DWBA</td>
<td>–</td>
<td>DWBA input is being entered</td>
</tr>
<tr>
<td>ECHO</td>
<td>on</td>
<td>Input lines are printed in the output</td>
</tr>
<tr>
<td>ELASTIC</td>
<td>off</td>
<td>Compute cross section in stand-alone scattering</td>
</tr>
<tr>
<td>FIT</td>
<td>off</td>
<td>Do an optical model fit and specify the parameters to be varied</td>
</tr>
<tr>
<td>FITE</td>
<td>off</td>
<td>The bound state energy is matched to the potential</td>
</tr>
<tr>
<td>FITMODE</td>
<td>1</td>
<td>Meaning depends on optical model fitter in use</td>
</tr>
<tr>
<td>FITV</td>
<td>on</td>
<td>The depth of the real part of the W.S.well is matched to the bound state energy</td>
</tr>
<tr>
<td>FREE</td>
<td>–</td>
<td>Remove an object from the allocator</td>
</tr>
<tr>
<td>GRIDSETUP</td>
<td>off</td>
<td>Grid setup input is being entered</td>
</tr>
<tr>
<td>HEADER</td>
<td>–</td>
<td>Defines the header for subsequent pages</td>
</tr>
<tr>
<td>INCOMING</td>
<td>off</td>
<td>The incoming scattering state is being defined</td>
</tr>
<tr>
<td>KEEP</td>
<td>–</td>
<td>Keep an object for subsequent Speakeasy use</td>
</tr>
<tr>
<td>KEYWORDS</td>
<td>–</td>
<td>List all keywords and their present values</td>
</tr>
<tr>
<td>LABANGLES</td>
<td>off</td>
<td>Angle grid is in laboratory</td>
</tr>
<tr>
<td>LINEAR</td>
<td>off</td>
<td>Only use linear extrapolation in the bound state search</td>
</tr>
<tr>
<td>LISTKEYS</td>
<td>–</td>
<td>List all keywords</td>
</tr>
<tr>
<td>LMCHOL</td>
<td>on</td>
<td>Use the LMCHOL sum-of-squares minimizer with</td>
</tr>
<tr>
<td>MINIM</td>
<td>off</td>
<td>Use the Fletcher variable-metric fitter</td>
</tr>
<tr>
<td>NEWPAGE</td>
<td>–</td>
<td>Go to new page for output</td>
</tr>
<tr>
<td>NOCHECKASYMP</td>
<td>on</td>
<td>Do not check asymptotic form of scattering solution</td>
</tr>
<tr>
<td>NOECHO</td>
<td>off</td>
<td>Do not print input lines in the output</td>
</tr>
<tr>
<td>NOELASTIC</td>
<td>on</td>
<td>Do not compute cross sections in stand-alone scattering problems</td>
</tr>
</tbody>
</table>
### A.5. Sub-keywords for DATA

The following are keywords that may appear within the parentheses that follow the `DATA` keyword.

<table>
<thead>
<tr>
<th>Keyword</th>
<th>Default</th>
<th>Meaning</th>
</tr>
</thead>
<tbody>
<tr>
<td>NOSAVEB</td>
<td>on</td>
<td>Do not save $B(L_x, M_x)$ cross sections</td>
</tr>
<tr>
<td>NOSAVEMXCROSS</td>
<td>on</td>
<td>Do not save $(L_x, M_x)$ cross sections</td>
</tr>
<tr>
<td>NRNLIMS</td>
<td>–</td>
<td>Compute partial overlaps of two functions</td>
</tr>
<tr>
<td>NSCATALOG</td>
<td>–</td>
<td>List names of all objects in allocator</td>
</tr>
<tr>
<td>NSDUMP</td>
<td>–</td>
<td>Print an object in the allocator</td>
</tr>
<tr>
<td>NSSTATUS</td>
<td>–</td>
<td>Print allocator statistics</td>
</tr>
<tr>
<td>NUMRNUM</td>
<td>–</td>
<td>Compute overlap of two functions</td>
</tr>
<tr>
<td>NZRDWBA</td>
<td>off</td>
<td>Input is for a nonzero-range DWBA</td>
</tr>
<tr>
<td>OUTGOING</td>
<td>off</td>
<td>Input is for the outgoing scattering state</td>
</tr>
<tr>
<td>PARAMETERSET</td>
<td>none</td>
<td>Specify a group of keyword definitions</td>
</tr>
<tr>
<td>POWELL65</td>
<td>off</td>
<td>Use the Powell 1965 sum-of-squares minimizer</td>
</tr>
<tr>
<td>PRINT</td>
<td>10001</td>
<td>Controls amount of printing</td>
</tr>
<tr>
<td>PRINTFIT</td>
<td>–</td>
<td>Print present E and V</td>
</tr>
<tr>
<td>PRINTWAV</td>
<td>–</td>
<td>Print last computed wavefunction</td>
</tr>
<tr>
<td>PROJECTILE</td>
<td>off</td>
<td>Projectile B.S.definition is being input</td>
</tr>
<tr>
<td>QUADRATIC</td>
<td>on</td>
<td>Use parabolic extrapolation in B.S.search</td>
</tr>
<tr>
<td>QUAVER</td>
<td>off</td>
<td>Use the QUAVER quasi-Newton fitter</td>
</tr>
<tr>
<td>RADIALINT</td>
<td>off</td>
<td>Input is for radial integration stage</td>
</tr>
<tr>
<td>REALWAVE</td>
<td>off</td>
<td>Real scattering waves for real pots</td>
</tr>
<tr>
<td>RENAME</td>
<td>–</td>
<td>Change the name of an object in the allocator</td>
</tr>
<tr>
<td>RESET</td>
<td>–</td>
<td>Initialize for a new calculation – allocator is CLEARed and all keywords set to initial status</td>
</tr>
<tr>
<td>RETURN</td>
<td>–</td>
<td>Terminate PTOLEMY</td>
</tr>
<tr>
<td>ROCORD</td>
<td>off</td>
<td>Use the ROCORD generalized minimizer</td>
</tr>
<tr>
<td>SAVEHS</td>
<td>off</td>
<td>Form factor integrals will be saved</td>
</tr>
<tr>
<td>SAVEB</td>
<td>off</td>
<td>Save the $B(L_x, M_x, \text{angle})$ for KEEP</td>
</tr>
<tr>
<td>SAVEMXCROSS</td>
<td>off</td>
<td>Save $(L_x, M_x)$ cross sections for KEEP</td>
</tr>
<tr>
<td>SCATTERING</td>
<td>off</td>
<td>Input is for stand-alone scattering</td>
</tr>
<tr>
<td>SIZE</td>
<td>$50 \times 10^6$</td>
<td>Allocator size in bytes</td>
</tr>
<tr>
<td>SKIPASYMP</td>
<td>off</td>
<td>Do not use Whittaker function as asymptotic form of B.S.wavefunctions</td>
</tr>
<tr>
<td>TARGET</td>
<td>off</td>
<td>Input is for Target B.S.</td>
</tr>
<tr>
<td>TSO</td>
<td>none*</td>
<td>Operating in interactive mode – errors allow corrections and retries</td>
</tr>
<tr>
<td>UNDEFINE</td>
<td>–</td>
<td>Set a keyword to undefined status</td>
</tr>
<tr>
<td>USEHS</td>
<td>off</td>
<td>Use previously computed form factor integrals</td>
</tr>
<tr>
<td>WRITENS</td>
<td>–</td>
<td>Write an object onto Fortran file 15</td>
</tr>
<tr>
<td>WRITESTEP</td>
<td>none</td>
<td>Interval (fm) at which wavefunctions are to be tabulated</td>
</tr>
</tbody>
</table>
### APPENDIX A. LIST OF ALL KEYWORDS

<table>
<thead>
<tr>
<th>Keyword</th>
<th>Default</th>
<th>Meaning</th>
</tr>
</thead>
<tbody>
<tr>
<td>ANGLE</td>
<td>–</td>
<td>C. m. angles of the data are being entered</td>
</tr>
<tr>
<td>ANGLESHEFT</td>
<td>0</td>
<td>Amount (in degrees) by which the angles are to be shifted in the laboratory frame</td>
</tr>
<tr>
<td>CHANNEL</td>
<td>–</td>
<td>Elastic channel of following data</td>
</tr>
<tr>
<td>CMANGLE</td>
<td>–</td>
<td>C. m. angles of data are being entered</td>
</tr>
<tr>
<td>CMSIGMA</td>
<td>–</td>
<td>C. m. cross sections in mb are being entered</td>
</tr>
<tr>
<td>ELAB</td>
<td>previous value</td>
<td>Laboratory energy data</td>
</tr>
<tr>
<td>ERROR</td>
<td>–</td>
<td>Errors have same units as data</td>
</tr>
<tr>
<td>LABANGLE</td>
<td>–</td>
<td>Lab. angles are being entered</td>
</tr>
<tr>
<td>LABSIGMA</td>
<td>–</td>
<td>Lab. cross sections in mb are being entered</td>
</tr>
<tr>
<td>MBERROR</td>
<td>–</td>
<td>Data errors are in millibarns</td>
</tr>
<tr>
<td>PERCENTERR</td>
<td>–</td>
<td>Data errors are in percent</td>
</tr>
<tr>
<td>POLARIZATION</td>
<td>–</td>
<td>Presently this data is ignored</td>
</tr>
<tr>
<td>RENORMALIZA</td>
<td>1</td>
<td>A quantity that will multiply each experimental cross section</td>
</tr>
<tr>
<td>SIGMA</td>
<td>–</td>
<td>Data is cross section in mb in ANGLE frame</td>
</tr>
<tr>
<td>SIGMATORUTH</td>
<td>–</td>
<td>Data is a ratio of cross section to Rutherford cross section</td>
</tr>
<tr>
<td>WEIGHT</td>
<td>1</td>
<td>Each term in the chi-squared sum for the present data group is multiplied by the WEIGHT</td>
</tr>
</tbody>
</table>

*A value will be computed by the program if one is not given by the user.*
Appendix B

Sample Ptolemy Jobs

The following is the input for several sample PTOLEMY jobs. These jobs use standard PTOLEMY input sequences and, to reduce confusion, do not explore the alternative ways of defining a problem. The outputs of these examples are separately available from the authors of this report.

B.1 Optical Model Fit

The following is an example of a five-parameter optical model fit to data at two energies. The depth of the real part of the potential is given an energy dependence while the rest of the potential parameters have no energy dependence. All of the potential radii will be equal to each other throughout the search while the real and imaginary diffusenesses will be separately varied. The search is first made with computed differential cross sections of moderate precision and is then repeated with more accurate values. Finally the differential cross sections for the two energies are printed on a uniform angular grid.

HEADER: EXAMPLE 1 - A TWO-ENERGY OPTICAL MODEL FIT
CHANNEL: 16O + 208PB
FIT ( R0=RI0=RC0 A AI V VE )
PARAMETERSET EL1
R0 = 1.3 A = .5 AI = .5 V = 40, VE = -.2 VI = 15
DATA ( ELAB=104 ANGLE SIGMATORU PERCENTER
$ KOVAR ET AL. 016 ON PB208 AT 104 MEV
The second example is of inelastic excitation. The B(E2) value is used to determine both the Coulomb and nuclear deformation parameters. The optical potential in the incoming state will also be used in the outgoing state and for computing the effective interaction.

//EXAMPLE2 JOB (F12345,5,0,2),REGION=200K,CLASS=C
B.2. TRANSFER REACTION

The following is a typical DWBA transfer input. Note the use of PRINT=2 to cause the cancellations in the computed radial integrals to be printed. Although both bound states involve a spin-orbit force, it is not necessary to specify JP since it can be uniquely determined by PTOLEMY from the other known spins and the given value of L.

HEADER: EXAMPLE 3 - A SIMPLE TRANSFER CALCULATION
REACTION: 208PB(16O 15N)209BI(5/2- 2.84) ELAB = 104
PARAMETERSET PB100A
PRINT = 2
PROJECTILE
NODES = 0 L = 1
R0 = 1.20 A=.65 RCO = 1.20
VSO = 7
;
TARGET
NODES = 1 L = 3
R0 = 1.28 A=.76
VSO = 6 RSO0 = 1.09 ASD = .6
;
INCOMING
RCO = 1.3 R0 = 1.2802 A = .5975 RIO = 1.2962 AI = .5424
V = 20 VI = 15
;
OUTGOING
;
$ THIS FINAL SEMI-COLON STARTS THE DWBA
KEYWORDS
RETURN