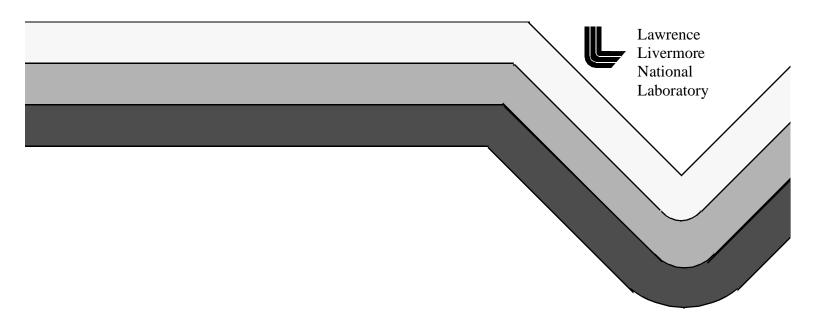
EQPT, A Data File Preprocessor for the EQ3/6 Software Package: User's Guide and Related Documentation (Version 7.0)

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December 17, 1992



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Preface

This report is the first in a set of documenting version 7.0 (version 3245.1090 under the old numbering system) of the EQ3/6 software package. This set includes:

I. The EQ3/6 Package Overview and Installation Guide (Wolery, 1992a).

II. The EQPT User's Guide (this report).

III. The EQ3NR Theoretical Manual and User's Guide (Wolery, 1992b).

IV. The EQ6 Theoretical Manual and User's Guide (Wolery and Daveler, 1992).

EQ3NR is the speciation-solubility code in the EQ3/6 package. EQ6 is a reaction path code and hence deals with the evolution of a water/rock system as reaction progress or time advances. EQPT is the EQ3/6 data file preprocessor.

The development of EQ3/6 has been supported by a number of programs concerned with geologic disposal of high level nuclear waste, including the Office of Nuclear Waste Isolation, the Salt Repository Project Office, the Waste Isolation Pilot Plant (through Sandia National Laboratory), the Nevada Nuclear Waste Storage Investigations, and the Yucca Mountain Site Characterization Project. Documentation for the package is aimed at satisfying the requirements of the U.S. Nuclear Regulatory Commission for software used for this purpose (Silling, 1983).

The Lawrence Livermore National Laboratory has not certified that EQ3/6 constitutes approved code for the conduct of quality affecting work for the Yucca Mountain Project.

No source codes or data files are reproduced in this report, nor are any computer media containing such items a part of this report or any of the other reports documenting this version of EQ3/6. The software itself must be obtained as described below.

The examples presented in this series of reports correspond to version 7.0 of the software and the R10 set of supporting thermodynamic data files. As of the date of publication of this report, the most recent version of the software is version 7.1 (containing bug fixes, but no enhancements), and the most recent set of data files is R16.

Agencies of the United States Government and their contractors may obtain copies of the software and its documentation from:

Energy Science and Technology Software Center P. O. Box 1020 Oak Ridge, TN 37831-1020

Telephone: (615) 576-2606

Requests to obtain the software under a licensing agreement should be addressed to:

Technology Transfer Initiatives Program, L-795 Attn: Diana (Cookie) West Lawrence Livermore National Laboratory P.O. Box 808 Livermore, CA 94550

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Contents

Glossary of Symbols

<i>a</i> , <i>a</i> '	Symbols used to represent cations in the notation of Harvie, Møller, and Weare (1984); see also X , X' .
a_i	Thermodynamic activity of the <i>i</i> -th aqueous solute species.
a_w	Thermodynamic activity of water.
$a_{\sigma\psi}$	Thermodynamic activity of the σ -th component of the ψ -th solid solution phase.
å	Generalized hard core diameter or "ion size" in aqueous solution.
å _i	Hard core diameter or "size" of the <i>i</i> -th aqueous solute species.
Α _{γ,10}	Debye-Hückel A parameter used in writing expressions for $log_{10} \gamma_i$.
A_{ϕ}	Debye-Hückel A parameter used in writing expressions for $ln a_w$.
b _{sr}	Stoichiometric reaction coefficient, the number of moles of the <i>s</i> -th aqueous species appearing in the <i>r</i> -th aqueous reaction; it is negative for reactants and positive for products.
b _{s\$}	Stoichiometric reaction coefficient, the number of moles of the <i>s</i> -th aqueous species appearing in the reaction for the dissolution of the ϕ -th pure mineral; it is negative for reactants and positive for products.
b _{sg}	Stoichiometric reaction coefficient, the number of moles of the s -th aqueous species appearing in the reaction for the dissolution of the g -th gas species; it is negative for reactants and positive for products.
b	A parameter theoretically equivalent to the product aB_{γ} and appearing in Pitzer's equations with an fixed value of 1.2.
B_{γ}	Debye-Hückel <i>B</i> parameter used in writing expressions for $ln \gamma_i$ or $log_{10} \gamma_i$.
C^{ϕ}_{MX}	Third order interaction coefficient for cation M and anion X .
e	The electron. In common thermodynamic formalism, this is usually a hypothetical species, not a real one.
Eh	Redox potential, volts. Theoretical equilibrium electrical potential of a redox couple; $Eh = \frac{2.303RT}{4F} (logf_{O_2} - 4pH - 2loga_w - logK_{\rm Eh}) \text{, where } f_{O_2} \text{ is understood to be the hypothetical equilibrium oxygen fugacity in aqueous solution.}$
f(I)	Debye-Hückel f function.
f(I)	Debye-Hückel f' function; $f(I) = df/dI$.
f_{O_2}	Oxygen fugacity.
F	The Faraday constant, 23062.3 cal/equiv-volt.
g	Subscript denoting a gas species.
g(x)	A function used to describe the ionic strength dependence of the second order interaction coefficient in Pitzer's equations.

Ι	Ionic strength.
Κ	Thermodynamic equilibrium constant.
K_{Eh}	Thermodynamic equilibrium constant for the half-reaction
	$2H_2O_{(1)} = O_{2(g)} + 4H^+ + 4e^-$
m _i	Molal concentration of the <i>i</i> -th aqueous solute species.
N, N'	Symbols denoting neutral species.
$N_{oldsymbol{\psi}}$	Site-mixing parameter for the ψ -th solid solution. If $N_{\psi} = 1$, the model is equivalent to a molecular- mixing model.
<i>O</i> ₂	Oxygen gas; in aqueous solution, this refers to a hypothetical species similar to e^{-} ; also symbolized as s_B .
$p_{k\psi}$	The <i>k</i> -th parameter used to compute the interaction coefficients W_{ψ} , $W_{i\psi}$, $W_{ij\psi}$, which in turn are used to compute the activity coefficients of end-member components in the ψ -th solid solution.
Р	Pressure, bars.
pe	Logarithm of the hypothetical electron activity; $pe = F Eh/(2.303 RT)$.
r	Subscript denoting an aqueous reaction.
R	The gas constant, 1.98726 cal/mol-°K.
S	Subscript denoting an aqueous species ($s = w$ implies $H_2O_{(l)}$).
<i>s</i> '	Subscript denoting s in the range from 1 to s_Q , excluding the cases $s = w$ and $s = s_B$.
<i>s</i> ''	Subscript implying the species formally associated with the aqueous reaction designated by r ($s'' = r + s_B$).
s _B	Subscript denoting the hypothetical aqueous species $O_{2(g)}$.
^S Q	The total number of aqueous master species; depending on the problem at hand, s_Q is equal to or greater than s_B .
s _T	Total number of aqueous species.
Т	Temperature, °K.
W	Subscript denoting water (e.g., a_{w} , the activity of water).
$W_{\psi} W_{i\psi} W_{ij\psi}$	
	Interaction coefficients used to compute the activity coefficients of end-member components in the ψ -th solid solution.
x	A general algebraic variable.
x_w	Mole fraction of water in aqueous solution.
x _{σψ}	Mole fraction of the σ -th end member of the ψ -th solid solution.
X. X'	Symbols denoting anions.

X, *X'* Symbols denoting anions.

 z_s Electrical charge of the *s*-th aqueous species.

- 2.303 Symbol for and approximation of *ln* 10. As an approximation, this is not in general sufficiently accurate approximation; this constant should be computed to full machine accuracy in a computer code in order to avoid both inaccuracy and inconsistency.
- α_1, α_2 Parameters appearing in Pitzer's equations.

 $\beta_{MX}^{(0)},\,\beta_{MX}^{(1)},\,\beta_{MX}^{(2)}$

	Observable second order interaction coefficient parameters for cation M and anion X .
γ_i	Molal activity coefficient of the <i>i</i> -th aqueous solute species.
8	Subscript denoting a chemical element.
ϵ_T	Total number of chemical elements in a system.
ζ_{NMX}	Observable third order interaction coefficient for neutral species N , cation M , and anion X .
$\theta_{MM'}$	Harvie, Møller, and Weare's (1984) notation for ${}^{S}\theta_{MM'}$.
${}^{S}\theta_{MM'}$	Observable short-range second order interaction coefficient for cations M and M' .
$\lambda_{ij}(I)$	Second-order interaction coefficient for the <i>i</i> -th and <i>j</i> -th aqueous solute species; in general, this is a function of the ionic strength.
$\lambda'_{ij}(I)$	The derivative of $\lambda_{ij}(I)$ with respect to ionic strength.
$\lambda_{MX}^{(0)},\lambda_{MX}^{(1)},$	$\lambda_{MX}^{(2)}$
	Second order interaction coefficient parameters for cation M and anion X .
λ_w	Second order interaction coefficient parameters for cation <i>M</i> and anion <i>X</i> . Rational (mole fraction) activity coefficient of water; $a_w = \lambda_w x_w$.
$λ_w$ λ _{σψ}	-
	Rational (mole fraction) activity coefficient of water; $a_w = \lambda_w x_w$.
$\lambda_{\sigma\psi}$	Rational (mole fraction) activity coefficient of water; $a_w = \lambda_w x_w$. Rational (mole fraction) activity coefficient of the σ -th end member of the ψ -th solid solution.
$\lambda_{\sigma\psi}$ $^{E}\lambda_{MM}(I)$	Rational (mole fraction) activity coefficient of water; $a_w = \lambda_w x_w$. Rational (mole fraction) activity coefficient of the σ -th end member of the ψ -th solid solution. The electrostatic part of $\lambda_{MM'}(I)$.
$λ_{σψ}$ ^E $λ_{MM}(I)$ ^S $λ_{MM'}$	Rational (mole fraction) activity coefficient of water; $a_w = \lambda_w x_w$. Rational (mole fraction) activity coefficient of the σ -th end member of the ψ -th solid solution. The electrostatic part of $\lambda_{MM'}(I)$. The short-range part of $\lambda_{MM'}(I)$; treated as a constant.
$λ_{σψ}$ ^E $λ_{MM'}(I)$ ^S $λ_{MM'}$ μ _{ijk}	Rational (mole fraction) activity coefficient of water; $a_w = \lambda_w x_w$. Rational (mole fraction) activity coefficient of the σ -th end member of the ψ -th solid solution. The electrostatic part of $\lambda_{MM'}(I)$. The short-range part of $\lambda_{MM'}(I)$; treated as a constant. Third-order interaction coefficient for the <i>i</i> -th, <i>j</i> -th, and <i>k</i> -th aqueous solute species.
$λ_{σψ}$ $E_{λ_{MM}}(I)$ $S_{λ_{MM'}}$ $μ_{ijk}$ $σ, σ'$	Rational (mole fraction) activity coefficient of water; $a_w = \lambda_w x_{w}$. Rational (mole fraction) activity coefficient of the σ -th end member of the ψ -th solid solution. The electrostatic part of $\lambda_{MM'}(I)$. The short-range part of $\lambda_{MM'}(I)$; treated as a constant. Third-order interaction coefficient for the <i>i</i> -th, <i>j</i> -th, and <i>k</i> -th aqueous solute species. Symbols denoting end-member components of a solid solution.
$λ_{σψ}$ $E_{λ_{MM'}}(I)$ $S_{λ_{MM'}}$ $μ_{ijk}$ $σ, σ'$ $σ_{T,ψ}$	Rational (mole fraction) activity coefficient of water; $a_w = \lambda_w x_{w}$. Rational (mole fraction) activity coefficient of the σ -th end member of the ψ -th solid solution. The electrostatic part of $\lambda_{MM'}(I)$. The short-range part of $\lambda_{MM'}(I)$; treated as a constant. Third-order interaction coefficient for the <i>i</i> -th, <i>j</i> -th, and <i>k</i> -th aqueous solute species. Symbols denoting end-member components of a solid solution. Total number of end members in the ψ -th solid solution.

EQPT, A Data File Preprocessor for the EQ3/6 Software Package: User's Guide and Related Documentation (Version 7.0)

Abstract

EQPT is a data file preprocessor for the EQ3/6 software package. EQ3/6 currently contains five primary data files, called data0 files. These files comprise alternative data sets. These data files contain both standard state and activity coefficient-related data. Three (com, sup, and nea) support the use of the Davies or B-dot equations for the activity coefficients; the other two (hmw and pit) support the use of Pitzer's (1973, 1975) equations. The temperature range of the thermodynamic data on these data files varies from 25°C only to 0-300°C. The principal modeling codes in EQ3/6, EQ3NR and EQ6, do not read a data0 file, however. Instead, these codes read an unformatted equivalent called a data1 file. EQPT writes a data1 file, using the corresponding data0 file as input. In processing a data0 file, EQPT checks the data for common errors, such as unbalanced reactions. It also conducts two kinds of data transformation. Interpolating polynomials are fit to data which are input on temperature grids. The coefficients of these polynomials are then written on the data1 file in place of the original temperature grids. A second transformation pertains only to data files tied to Pitzer's equations. The commonly reported observable Pitzer coefficient parameters are mapped into a set of *primitive* parameters by means of a set of conventional relations. These primitive form parameters are then written onto the **data1** file in place of their observable counterparts. Usage of the primitive form parameters makes it easier to evaluate Pitzer's equations in EQ3NR and EQ6. EQPT and the other codes in the EQ3/6 package are written in FORTRAN 77 and have been developed to run under the UNIX operating system on computers ranging from workstations to supercomputers.

1. Introduction

EQPT is a data file preprocessor. It is part of the EQ3/6 software package (see Wolery, 1992a). It replaces the EQTL code (see Wolery, 1983). This report describes EQPT in version 7.0 (version 3245.1090 in the old numbering system) of this package (see the EQ3/6 Package Overview and Installation Guide, Wolery, 1992a). Other codes in the package include EQ3NR (Wolery, 1992b), a speciation-solubility code, and EQ6 (Wolery and Daveler, 1992), a reaction path code. The relationship of EQPT code to EQ3NR, EQ6, and the set of supporting thermodynamic data files is shown in Figure 1. This figure depicts the flow of information involving these codes. At present, there are five distinct data files, denoted by the suffixes **com**, **sup**, **nea**, **hmw**, and **pit**. These are provided in formatted ASCII and are called **data0** files. The full name of such a file ordinarily has a form exemplified by **data0.com.R10**, where the **R10** is a stage number (a configuration control identifier). On some systems, it is necessary to compress this to a form exemplified by **da0com.R10** (see Wolery, 1992a).

The user of EQ3NR or EQ6 must select which of the five data files is most appropriate to a given problem. Each data file corresponds to a general formalism for treating the activity coefficients of aqueous species and contains the relevant activity coefficient data as well as standard state thermodynamic data (e.g., dissociation constants). The **com**, **sup**, and **nea** data files are specific to a general extended Debye-Hückel formalism and can be used by EQ3NR and EQ6 with either the Davies (1962) equation or the B-dot equation (Helgeson, 1969). These equations are only

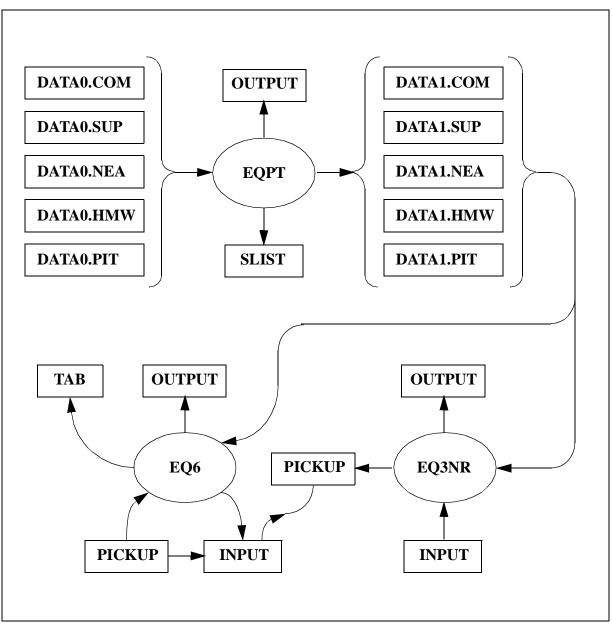


Figure 1. The flow of information among the computer codes EQPT, EQ3NR, and EQ6. Computer codes are represented by ovals, files by rectangles.

valid in relatively dilute solutions. The **hmw** and **pit** data files are specific to the formalism proposed by Pitzer (1973, 1975) and can be used to model solutions extending to high concentrations. However, the scope of chemical components covered is smaller. The temperature limits on the data files also vary, from 25°C only to 0-300°C.

Some important data file characteristics are given in Table 1. The **com** (for "composite") data file is the largest of the three data files specific to the extended Debye-Hückel formalism. It is a product of Lawrence Livermore National Laboratory (LLNL) drawing on many data sources, includ-

ing those on which the other four data files are based. The **sup** data file is based entirely on SUPCRT92 (Johnson, Oelkers, and Helgeson, 1992), a software package and thermodynamic data base founded on the work of Helgeson and Kirkham (1974ab, 1976), Helgeson et al. (1978), Tanger and Helgeson (1988), Shock and Helgeson (1988, 1989, 1990), Shock, Helgeson, and Sverjensky (1989), Johnson and Norton (1991), and Shock et al. (1992). The **nea** data file is based entirely on Grenthe et al. (1989, draft report), a product of the Data Bank of the Nuclear Energy Agency of the European Community. This work has recently been published as Grenthe et al. (1992). The **hmw** data file is based on Harvie, Møller, and Weare (1984). The **pit** data file is based mostly on data summarized by Pitzer (1979). All five data files are maintained at LLNL in a relational data base described by Delany and Lundeen (1991). This relational data base is part of the Yucca Mountain Site Characterization Project's Technical Data Base.

The **sup** data file has a high level of internal consistency among the standard state thermodynamic data. In addition, the temperature-pressure dependence of these data are represented by a suite of equations of state for minerals, gases, and aqueous species that are well established in the geochemical literature (see references noted above). This data file covers a wide range of chemical elements and species of interest in the study of rock/water interactions (e.g., components which make up the major rock-forming and ore-forming minerals). It also includes a large number of organic species, mostly of small carbon number (C_2 - C_8). The **nea** data file is something of a specialty item. Its strongest point is a thorough representation of the thermodynamics of uranium species.

The **com** (composite) data file encompasses a much broader range of chemical elements and species. It includes the data found on the **sup** and **nea** data files, with preference given to data from the former in cases of overlap. It also includes some data found in the **hmw** data file, as well as other data which do not appear in any of the other data files. Some of these data are estimates based on correlations or extrapolations (as to higher temperature), and are not tied directly to experimental measurements. The **com** data file thus represents a melange of data, which by its nature offers less assurance of internal consistency. However, this offers the only means presently available for modeling aqueous solutions with a high degree of compositional complexity, such as the fluids expected to be found in and about a facility for the geologic disposal of industrial or nuclear waste (e.g., the potential repository for high-level nuclear waste at Yucca Mountain, Nevada).

The **hmw** data file has the highest degree of internal consistency of any of the five data files, including mutual consistency of activity coefficient data and standard state thermodynamic data. It can be applied to dilute waters or concentrated brines. However, it only treats the set of components present in the "sea-salt" system (the major cations and anions present in seawater, including carbonate and bicarbonate). The geochemically important components aluminum and silica are not included. Also, this data file is limited to a temperature of 25°C. The **pit** data file can also be applied to concentrated brines. It covers a larger set of components, but these mostly involve other cations and anions of strong electrolytes. Examples include lithium and bromide. This data file nominally covers the temperature range of 0-100°C. However, it represents a melange of data, not a carefully crafted internally consistent set.

What one can do with EQ3NR or EQ6 depends to a high degree on what is on the supporting data file and if so how the species and reactions on the data file are organized. For example, iron does

File Name (Suffix)	Source	Activity Coefficient Formalism	Temperature Limits	Number of Chemical Elements	Number of Basis Species	Number of Aqueous Species	Number of Pure Minerals	Number of Solid Solutions	Number of Gas Species
com	GEMBOCHS (LLNL)	Extended Debye- Hückel	0-300°C	78	147	852	886	12	76
sup	SUPCRT92	Extended Debye- Hückel	0-300°C	69	105	315	130	0	16
nea	NEA draft report	Extended Debye- Hückel	0-300°C	32	50	158	188	0	76
hmw	Harvie, Møller, and Weare (1984)	Pitzer's Equations	25°C only	9	13	17	51	0	3
pit	Pitzer (1979)	Pitzer's Equations	0-100°C	52	62	68	381	0	38

not appear on the **hmw** data file, so this file can not be used to model the behavior of iron in brinemineral systems. Similarly, uranium does not appear on the **sup** data file. Even if a chemical element does appear on a given data file, the corresponding species required for a given problem must also appear on it.

EQPT processes these **data0** files one at a time (looking for a file named simply **data0**, though these files are normally stored under names which include the relevant suffixes) and writes a corresponding unformatted data file, which is called simply **data1**. These are also normally stored under names including the relevant suffixes (e.g., **data1.com.R10**). To run EQ3NR or EQ6, the user must provide one of these files, which is known to each code simply as **data1**.

To process all five data files running EQPT directly thus requires the user to do a lot of renaming of the various files. This is not very convenient. The export package includes a UNIX shell script called **runeqpt** to make the job easier. This script and its usage are described in the EQ3/6 Package Overview and Installation Manual (Wolery, 1992a). Caution: this script may require local modification, as it needs data for the location within the local directory structure of both the **data0** files and the EQPT executable file The script is invoked by commands of the form:

runeqpt R10 all

(processes all data0 files with stage number R10) or:

runeqpt R10 com

(processes only the data file data0.com.R10).

The **runeqpt** script renames all of the files produced, incorporating the data file key and stage number of each **data0** file processed. For example, the **data1** file for **data0.com.R10** will be named **data1.com.R10**, and the **slist** file will be named **slist.com.R10**. Naming the **data1** files in this manner facilitates running EQ3NR and EQ6 under the shell script **runeq36** (se the EQ3/6 Package Overview and Installation Manual, Wolery, 1992a).

Note that EQPT in its present form has no **input** file and no user options. In porting EQ3/6, the user need only run EQPT on each of the **data0** files provided. This should be a simple process. The actual purpose of this manual is to comply with the NUREG documentation requirements (Silling, 1983) and to provide information concerning the data file structure and its processing that might be useful to users who modify the original data files or make up data files of their own.

The data file preprocessor EQPT performs a number of functions. It checks the composition, charge, and reaction coefficient data on a **data0** file for internal consistency and fits interpolating polynomials to various temperature dependent data which are organized on the **data0** file on temperature grids. Such data include certain activity coefficient parameters, such as Debye-Hückel $A_{\gamma,10}$ and B_{γ} , and the equilibrium constants for the reactions represented on the data file. In addition, in the case of data files specific to the formalism of Pitzer's equations, observable interaction coefficients are mapped to a set of conventionally defined primitive interaction coefficients. EQPT then writes the **data1** file corresponding to the input **data0** file. It also writes a formatted equivalent, called **data1f**. The user may examine this if desired, but this file is not otherwise used for any purpose. In addition, EQPT writes to a screen file and an **output** file, both of which are

generally significant only if an error condition is encountered. In addition, it writes an **slist** (species list) file. This is very useful to the user, as it lists the species that are represented on the data file and identifies which species are in the strict and auxiliary basis sets. If no errors are encountered, the **output** and **slist** files are nearly identical.

When processing data files corresponding to the Pitzer formalism, EQPT also writes two files called **dpt1** and **dpt2**. The former contains the original, observable Pitzer coefficient data. The latter contains the equivalent conventional primitive Pitzer coefficient data. These files are vestigial and may be discarded.

Chapter 2 describes the organization of species and reactions. This is centered on the concept of a set of basis species, including the concept of an auxiliary basis set. Chapter 3 describes the types of thermodynamic data present on the data files, how they are represented, and how they are transformed or mapped by EQPT when it writes the **data1** file. Chapter 4 describes the structure of the **data0** files and the corresponding **data1** files. Chapter 5 describes the code architecture. Appendix A contains a glossary of the major code variables. The source code modules are listed and briefly described in Appendix B (for a similar treatment of EQLIB modules, see Appendix B of the EQ3/6 Package Overview and Installation Guide, Wolery, 1992a). Appendix C contains a list of error messages generated by EQPT modules, along with related notes (see Appendix C of Wolery, 1992a, for a similar list for EQLIB modules). Appendix D contains notes pertaining to known bugs and such. Appendix E contains the **slist** files for the **com** and **hmw** data files.

EQPT and the other codes in the EQ3/6 software package are written in FORTRAN 77 and have been developed to run under UNIX operating systems on computers ranging from workstations to supercomputers, including Sun SPARCstations, VAXes (ULTRIX operating system), Alliants (CONCENTRIX operating system), and Crays (UNICOS operating system). They are fairly readily portable to VAX computers running the non-UNIX VMS operating system. They may be portable as well to 386 and 486 PCs. Platforms used at LLNL include Sun SPARCstations and an Alliant FX/80. For details concerning platforms, see the EQ3/6 Package Overview and Installation Guide (Wolery, 1992a).

2. Organization of Species and Reactions

The concept of a set of basis species is critical to the data base organization. EQ3/6 further utilizes the concept of a set of auxiliary basis species. The purpose of the present chapter is to present a brief description of these concepts as they pertain to the structure of the EQ3/6 data file. These concepts are discussed in more detail and from the viewpoint of the code user in Chapter 5 of the EQ3NR Theoretical Manual and User's Guide (Wolery, 1992b).

A *basis* species is a species used as a general "building block" for writing chemical reactions in a standardized format that is convenient for chemical modeling. Each species appearing in a reaction is a basis species, except for one non-basis species which is associated with the reaction itself. In the format used in EQ3/6, each reaction always destroys this associated non-basis species. This is illustrated by the reaction:

$$CaHCO_3^+ = Ca^{2+} + HCO_3^- \tag{1}$$

where the ion pair $CaHCO_3^+$ is the associated (non-basis) species and the other three species appearing in the reaction are basis species. Because it is a non-basis species, $CaHCO_3^+$ does not appear in any other reaction on the data file.

Each basis species on an EQ3/6 data file must be an aqueous species. In addition, it should usually be composed of no more than three chemical elements, no more than one of which may be other than oxygen or hydrogen. Thus, $H_2O_{(l)}$, H^+ , Na^+ , Ca^{2+} , Cl^- , UO_2^{2+} , SO_4^{2-} , and HCO_3^- are examples of species which can be and are used as basis species. Species such as $CaHCO_3^+$ and $UO_2CO_{3(aq)}$ should not appear as basis species on the data file, because they do not meet this requirement. However, they can be switched into the active basis set via basis switching options available in EQ3NR and EQ6 (See Wolery, 1992b, and Wolery and Daveler, 1992). These restrictions on basis species composition are employed in order to insure that the corresponding mass balance totals for an aqueous solution pertain to physically measurable quantities (see Chapter 5 of the EQ3NR Theoretical Manual and User's Guide, Wolery, 1992b). These are not hard and fast rules. An exception will be noted later in this chapter.

Each *strict* basis species is associated one-to-one with a a chemical element, with the exception of one which is associated with a redox parameter. The strict basis species used to write oxidation-reduction reactions in EQ3/6 is O_2 , which is treated as a fictive aqueous species. The associated redox parameter is the oxygen fugacity. A strict basis set is a minimal basis set; it represents the smallest possible basis set for a given set of chemical components, and corresponds to the case of complete chemical equilibrium within the aqueous solution. A strict basis species has no associated reaction.

It is advantageous to allow for an *auxiliary* basis set. Its members are usually similar species, most often representing a chemical element in a different oxidation state. An auxiliary basis set permits consideration of disequilibrium between a species in this set and a related species in the strict basis set. They are related by a reaction that is associated with each member of the auxiliary

basis set. For example, if Fe^{2+} is in the strict basis set and Fe^{3+} is in the auxiliary basis set, the reaction may be written as:

$$Fe^{3+} + \frac{1}{2}H_2O = Fe^{2+} + H^+ + \frac{1}{2}O_2$$
 (2)

The auxiliary basis set is preserved in its original form by EQPT. In EQ3NR or EQ6, an auxiliary basis species may be eliminated from the *active* basis set by using the associated reaction to rewrite all other reactions originally written in terms of that species so that it no longer appears in them. For example, the reaction:

$$FeCl_3^o = Fe^{3+} + 3Cl^{-}$$
 (3)

is combined with eq (2) to obtain:

$$FeCl_{3}^{o} + \frac{1}{2}H_{2}O = Fe^{2+} + H^{+} + \frac{1}{2}O_{2} + 3Cl^{-}$$
(4)

The ferric trichloride complex then appears to be a complex of the ferrous ion. The mass action equation corresponding to reaction (2) is activated as a constraint, so the ferric ion itself (in addition to its complexes) is treated as a "complex" of the ferrous ion.

In a geochemical modeling code such as EQ3NR or EQ6, the incorporation of an auxiliary basis set allows the code user to treat a species in this set as either a basis species or as a non-basis (dependent) species. If it is treated as a basis species, an additional mass balance relation is defined and the reaction relating the species to a corresponding basis species is allowed to be in disequilibrium (the degree of which can be calculated). Otherwise, this auxiliary basis species (and all of its dependent species) are treated as dependent species of that corresponding basis species. They are then folded into the mass balance for that species.

Each reaction is associated with a non-basis or auxiliary basis species, and each non-basis or auxiliary basis species has an associated reaction. If an auxiliary basis species is to be treated as a basis species in EQ3NR or EQ6, its associated reaction is ignored, except for the purpose of computing the degree of disequilibrium. If an auxiliary basis species is eliminated from the active basis set, it is treated as a non-basis species, and its reaction is utilized in the same manner as that of a non-basis species. Thus, one should place a species in the auxiliary basis set if one wishes to allow the possibility of treating the corresponding reaction in a state of disequilibrium.

As a general rule, reactions of non-basis on an EQ3/6 data file should be written so that the associated species is transformed into the chemically most similar basis species. This means that one should attempt to preserve such things as oxidation states or molecular moieties (structural groups), on the theory that the associated reactions are more likely to be in a state of equilibrium. For example, one would write on the data file reaction (3) for the ferric trichloride complex (yielding the ferric ion). If one wrote instead reaction (2) (yielding the ferrous ion), this complex would always be treated as falling under the mass balance for the ferrous ion. This would be erroneous in the case in which ferric-ferrous disequilibrium is assumed and a separate mass balance is employed for the ferric ion. It is not always possible to satisfy this guideline. The currently existing data files include ferrous and ferric ions as basis species, but not a dissolved iron species in the zero oxidation state. Metallic iron (Fe) requires a reaction, but there is no corresponding basis species on these files in the zero oxidation state. The reaction is therefore written instead with the minimal amount of oxidation-reduction:

$$iron + \frac{1}{2}O_2 + 2H^+ = Fe^{2+} + H_2O_{(l)}$$
(5)

Note that if the data file did contain a dissolved iron species in the zero oxidation state, such a species should be placed in the basis set.

A number of organic species appear in the **sup** and **com** data files. HCO_3^- is the strict basis species corresponding to the element carbon. The following carbon-bearing species appear on these data files as auxiliary basis species:

- ortho-phthalate ('o-(phth)--'; **com** file only)
- acetic acid ('acetic acid(aq)')
- acetone(aq)
- benzene(aq)
- ethane(aq)
- glycine(aq)
- methanamine(aq)
- methane(aq)
- methanol(aq)

Each of these except dissolved methane $(CH_{4(aq)})$ is treated as the parent of other organic species. For example, acetic acid is $(CH_3COOH_{(aq)})$ taken as the parent of propanoic acid $(CH_3CH_2COOH_{(aq)})$, the reaction for the latter being:

propanoic
$$acid(aq) + \frac{1}{2}O_2 = \frac{3}{2}acetic \ acid(aq)$$
 (6)

Here there is an attempt to preserve organic moieties (in this case, the -*COOH* group). Similarly, glycine (the simplest amino acid, $CH_2NH_3COOH_{(aq)}$) is treated as the parent of the other amino acids. Note that in dealing with organics, it is nearly impossible to avoid oxidation-reduction in writing reactions for the non-basis organic species. Also, methanamine ($CH_2NH_{3(aq)}$) violates the usual compositional guideline in that it is composed of both carbon and nitrogen.

The associated reaction for each of the above organic auxiliary basis species is written on the data files in the R7 and R10 sets so that the species is oxidized to bicarbonate. On data files in the R16 and higher sets, the species 'acetic acid(aq)' plays the role of parent to the others, with the exception of aqueous methane (i.e., their associated reactions yield acetic acid, not bicarbonate). Acetic acid then functions as a master organic species; aqueous methane is treated as "inorganic," an admittedly arbitrary choice. To illustrate the result, the user of EQ3NR can then avoid all organics in a calculation simply by specifying a zero concentration of acetic acid on the **input** file. To avoid aqueous methane as well, it must also be treated in this manner. Using data files from the earlier sets, to avoid all organics one must do this for each of the organic auxiliary basis species, and for aqueous methane as well, if it is not desired.

In the **sup** and **com** data files in the R16 set, the auxiliary basis species $S_2^{2^-}$ is similarly treated as the master polysulfide species (other polysulfides including $S_3^{2^-}$, $S_4^{2^-}$, and $S_5^{2^-}$). Also, the species $S_2O_3^{2^-}$ is treated as the master species for related partially oxidized sulfur species, including $S_2O_4^{2^-}$, $S_2O_5^{2^-}$, $S_2O_6^{2^-}$, and $S_2O_8^{2^-}$. Again, the rationale is to make it easy for the user to eliminate such species in model calculations when it is so desired.

All non-aqueous species (pure minerals, end-member components of solid solutions, non-aqueous liquids, and gases) are treated on the data files as non-basis species. Thus, all reactions for such species take the form of dissolution reactions.

3. Thermodynamic Data: Representations, Transformations, and Mappings

The purpose of this chapter is to note the types of thermodynamic data that appear on an EQ3/6 data file, and to discuss certain transformations and mappings made from these data by EQPT. The purpose of thermodynamic data is to allow evaluations of mass action equations. This is facilitated by the representations, transformations, and mappings described below.

There are fundamentally two different categories of thermodynamic data. This can be illustrated by the reaction for the dissolution of halite (*NaCl*):

$$NaCl = Na^{+} + Cl^{-}$$
⁽⁷⁾

The corresponding mass action equation can be written as:

$$\log K_{NaCl} = \log m_{Na^+} + \log \gamma_{Na^+} + \log m_{Cl} + \log \gamma_{Cl}$$
(8)

where *K* denotes the equilibrium constant, *m* the molality, and γ the molal activity coefficient. The equilibrium constant is an example of *standard state* thermodynamic data. The activity coefficients are examples of *excess* thermodynamic data. Although referred to as "data," these entities are perhaps more properly referred to as functions. They may in fact be calculated from other "data."There are other examples of kinds of thermodynamic data in each category, representing the temperature and pressure derivatives of these functions. For example, standard partial molar volumes are standard state data. However, these other kinds of thermodynamic data are not directly used in version 7 of EQ3/6.

Most "thermodynamic data bases" emphasize the standard state kind of data. However, the two kinds of data are closely linked. For example, reaction (8) allows computation of the solubility of halite in aqueous solutions. Conversely, measurements of halite solubility can be used to obtain the equilibrium constant, but not independently of the activity coefficients. In fact, one could (and in the most rigorous sense should) use such measurements to obtain both the equilibrium constants and the activity coefficients. This is true, despite the fact that the activity coefficients are obtainable in principle solely from measurements of the osmotic coefficient (cf. Pitzer, 1973).

In principle, a thermodynamic data base is internally consistent only if the combined set of standard state and excess thermodynamic data are mutually consistent. This has been achieved quite rarely; the work of Harvie, Møller, and Weare (1984) provides one of the few examples. It is difficult to provide a high level of internal consistency even among the standard state thermodynamic data (for an example of a large data base with such consistency, see the SUPCRT92 of Johnson, Oelkers, and Helgeson, 1992). To deal with the issue of mutual consistency among standard state and excess thermodynamic data, EQ3/6 data files are designed to contain both types on a given file. This by itself only permits mutual consistency; it does not guarantee it.

In the remainder of this chapter, we discuss first the standard state thermodynamic data, then the excess thermodynamic data. Representations and any transformations or mappings are described for each member of each category of data.

3.1. Standard State Thermodynamic Data

The only example of standard state thermodynamic data used in version 7 of EQ3/6 is the equilibrium constant, taken as log *K*. This is represented as a sequence of values on a temperature grid (0-25-60-100-150-200-250-300°C; see Chapter 4). The equilibrium constant is a function of temperature as well as pressure. However, the pressure is currently taken as a function of the temperature, with values defined on the same temperature grid. The pressure is 1.013 bar up to 100°C, and follows the steam/liquid water equilibrium pressure at higher temperature. Thus, log K(T) is really treated as log K(T,P(T)). EQ3/6 calculations are presently restricted to this pressure curve.

There is a log K grid for each reaction. There is one reaction for each aqueous species not in the strict basis set, one for each pure mineral, and one for each gas species. In addition, there is an equilibrium constant (log K_{Eh}) for the following special redox reaction:

$$2H_2O_{(l)} = O_{2(g)} + 4H^+ + 4e^-$$
(9)

where *e*- is the fictive aqueous electron. This reaction is built into EQ3/6. It is used to compute secondary redox variables (the redox potential *Eh* and the electron activity function *pe*) from the primary redox variable, the oxygen fugacity (f_{O_2}) .

The grid representation is inconvenient for modeling code calculations, because it may be desired to make calculations for temperatures not corresponding to one of the grid points. EQPT transforms the gridded data, replacing it with a set of coefficients for interpolating polynomials. One interpolating polynomial is applied to the 0-25-60-100°C part, another to the 100-150-200-250-300°C. This division respects the discontinuity of the temperature dependence of the grid pressure. It also assures a polynomial of only moderate order.

For a detailed discussion of the subject of fitting interpolating polynomials, the reader is referred to Chapter 1 of Carnahan, Luther, and Wilkes (1969) (or the appropriate section of almost any introductory text dealing with numerical methods). The interpolating polynomials used by EQPT are particularly simple in that they are exact; that is, they pass through all of the data points used in the fitting (a given fitting is limited to a corresponding temperature range). The 0-25-60-100°C part of the grid offers at most four data points, the 100-150-200-250-300°C part, at most five. Thus, the maximum order of the interpolating polynomial is three and four, respectively. Continuity at 100°C is guaranteed if a valid value for this temperature is present on the data grid.

The actual order of the interpolating polynomial depends on the number of valid points. Some points on the grid may be empty due to lack of data. The lack of data condition is marked in the case of log *K* grids by entering a value of "500.0000." Only valid points are used in the fitting. If only a 25°C value is present in the lower temperature part of the grid, the code fits a zero-th order polynomial (i.e., a constant). In order words, the 25°C value is extrapolated over the entire range. If there are no valid points in a given range, the code fits a zero-th order polynomial with a value of "500.0000." The effect of this is to suppress the associated species.

3.2. Excess Thermodynamic Data

The only kind of excess thermodynamic "data" used in the present version of EQ3/6 is the activity coefficient. This is actually treated as a function of other, related data. We will first consider the activity coefficients of aqueous species, then the activity coefficients of components of solid solution phases. The present version of EQ3/6 does not address non-aqueous liquid phases (e.g., a mixture of liquid hydrocarbons), hence it has no provision for treating the activity coefficients of component species of such phases. Nor does it address a gas phase, which would require the consideration of fugacity coefficients.

3.2.1. Activity Coefficients of Aqueous Species

The present version of EQ3/6 offers two formalisms for treating the activity coefficients of aqueous species (for a detailed discussion, see Chapter 3 of the EQ3/6 Theoretical Manual and User's Guide, Wolery, 1992b). The first of these includes simple extensions of the standard Debye-Hückel model, and consists of options for the Davies (1962) equation and the B-dot equation (Helgeson, 1969). The second is based on Pitzer's (1973, 1975, 1979, 1987) equations. The data requirements for the two kinds of formalisms are quite different and will be discussed below. The **com**, **nea**, and **sup** data files correspond to the first formalism, the **hmw** and **pit** data files, to the second.

In the discussion which follows, we give the equation for the solute activity coefficient (γ_i) as a means of introducing the parameters. A complete model for activity coefficients in aqueous solution also requires a corresponding equation for the activity of water (a_w) or its mole fraction activity coefficient (λ_w). These equations are presented in Chapter 3 of Wolery (1992b). They introduce no additional parameters (thermodynamic consistency requires that they do not), hence are not reproduced here.

3.2.1.1. Extended Debye-Hückel Formalism

The extended Debye-Hückel formalism is represented in the present version of EQ3/6 by the

Davies (1962) and the B-dot (\dot{B}) equation of Helgeson (1969). These are sufficiently accurate for geochemical applications only in relatively dilute solutions (having ionic strengths of at most 1 molal). Their chief advantage is that the data requirements posed by these models are quite minimal.

3.2.1.1.1. The Davies (1962) equation

The Davies (1962) equation is:

$$\log \gamma_i = -A_{\gamma, 10} z_i^2 \left(\frac{\sqrt{I}}{1 + \sqrt{I}} + 0.2I \right) \tag{10}$$

This is a simple extended Debye-Hückel model, to which it reduces if the "0.2*I*" part is removed. The only data required is for the Debye-Hückel $A_{\gamma, 10}$ parameter. This is here written with a subscript "10" to note consistency with the base ten logarithm on the left hand side of eq (10). The only species-specific data required is for the electrical charge, which is actually an intrinsic parameter.

The use of this option requires a supporting data file consistent with the use of a simple extended Debye-Hückel model (e.g., **com**, **nea**, or **sup**). The parameter $A_{\gamma, 10}$ is represented on the data file by a data grid completely analogous to those used to represent log *K* values. EQPT transforms this data grid into interpolating polynomials in the same manner.

3.2.1.1.2. The B-dot equation

The B-dot equation (Helgeson, 1969) is:

$$log\gamma_{i} = -\frac{A_{\gamma,10}z_{i}^{2}\sqrt{I}}{1 + \mathring{a}_{i}B_{\gamma}\sqrt{I}} + \dot{B}I$$
⁽¹¹⁾

Here $A_{\gamma,10}$ is the Debye-Hückel A parameter discussed above, B_{γ} is the Debye-Hückel *B* parameter, \dot{B} is the B-dot parameter, and \dot{a}_i is the hard core diameter of the species.

The use of this option requires a supporting data file consistent with the use of a simple extended Debye-Hückel model (e.g., **com**, **nea**, or **sup**). The parameters $A_{\gamma,10}$, B_{γ} , and \dot{B} are represented on the data file by data grids completely analogous to those used to represent log *K* values. EQPT transforms these data grids into interpolating polynomials in the same manner. The hard core diameters (a_i) are species-specific. They are taken to be constants. Values are assigned in a section of the data file described in Chapter 4.

When the B-dot option is chosen, the B-dot equation itself is only applied to charged species. Other equations are actually used for uncharged species. Each species is also assigned an **insgfl** flag; this flag is ignored for charged species. It appears in the same section of the data file as the hard core diameter (see Chapter 4).

For dissolved gases and other neutral species not of a strongly polar nature, the practice is to assign the value of the activity coefficient of aqueous CO_2 in otherwise pure sodium chloride solutions of the same ionic strength (Garrels and Thompson, 1962; Helgeson, 1969). This is computed from the following expression after Drummond (1981, p. 19):

$$ln\gamma_i = \left(C + FT + \frac{G}{T}\right)I - (E + HT)\left(\frac{I}{I+1}\right)$$
(12)

(see Chapter 3 of Wolery, 1992b) where *T* is the absolute temperature and C = -1.0312, F = 0.0012806, G = 255.9, E = 0.4445, and H = -0.001606. The coefficients appearing in eq (12) appear directly on the data file, and are not transformed or mapped in any way by EQPT. This treatment is marked by an **insgfl** value of 0. Note that $\log \gamma_i$ is computed from $\ln \gamma_i$

Following the recommendation of Garrels and Christ (1965, p. 70), the activity coefficients neutral aqueous species of a polar nature are set to unity; i.e., the equation is:

$$\log \gamma_i = 0 \tag{13}$$

This requires no additional data. This treatment is marked by an **insgfl** value of -1.

3.2.1.2. Pitzer's Equations

Pitzer's (1973, 1975, 1979, 1987) equations are based on a virial expansion theory and require a fairly large set of interaction coefficients to model systems of much complexity. Apart from that, they require one Debye-Hückel parameter, which is discussed below. The equations themselves may be written in more than one form, depending on the type of interaction coefficients one wishes to use. It is customary to report interaction coefficients of the *observable* type. Therefore, this type of interaction coefficient appears on the corresponding EQ3/6 data file (e.g., **hmw** or **pit**). However, EQ3/6 actually calculates the activity coefficients using interaction coefficients of the *primitive* type. EQPT maps observable interaction coefficients to a conventionally defined set of primitive interaction coefficients.

In the original theoretical form (Pitzer, 1973), the activity coefficient was written in term of the primitive interaction coefficients λ_{ii} and μ_{iik} :

$$\ln \gamma_i = \left(\frac{z_i^2}{2}\right) f(I) + 2\sum_i \lambda_{ij}(I)m_j + \sum_{ik} \left(\left(\frac{z_i^2}{2}\right)\lambda'_{jk}(I) + 3\mu_{ijk}\right)m_j m_k$$
(14)

Here f(I) is a Debye-Hückel function, f(I) is its derivative df/dI, the λ_{ij} are second order interaction coefficients, $\lambda'_{ij}(I)$ is the derivative $d\lambda_{ij}/dI$, and the μ_{ijk} are third order interaction coefficients. As is implied, the λ_{ij} are treated as functions of the ionic strength. The sums in the interaction coefficient terms are actually double and triple sums. This is the form actually evaluated by EQ3NR and EQ6. Its main virtue is that it is simple and compact, but completely general. It applies to any mixture of cations, anions, and uncharged solute species

The Debye-Hückel model used in Pitzer's equations is not the usual Debye-Hückel-charging formulation exemplified in the Davies or B-dot equations, but a different one derived by Pitzer (1973) and called the Debye-Hückel-osmotic model. The function f(I) is given by:

$$f(I) = -\left(\frac{4A_{\phi}I}{b}\right)ln\left(1 + b\sqrt{I}\right)$$
(15)

Here A_{ϕ} is a Debye-Hückel parameter is related to the more familiar $A_{\gamma,10}$ by:

$$A_{\phi} = \frac{2.303A_{\gamma, 10}}{3} \tag{16}$$

The parameter *b* is assigned a constant value of 1.2 (Pitzer, 1973). Different values have been used for the A_{ϕ} parameter at 25°C. It is important to use the value 0.392 with the Harvie, Møller, and Weare (1984) model of the "sea salt" system, not the stated value of 0.39 (see Plummer et al., 1988, p. 3).

Following Pitzer (1973) and Pitzer and Mayorga (1974) (see Chapter 3 of Wolery, 1992b), the second order interaction coefficient for cation-anion interactions is described by:

$$\lambda_{MX}(I) = \lambda_{MX}^{(0)} + \lambda_{MX}^{(1)}g(\alpha_1\sqrt{I}) + \lambda_{MX}^{(2)}g(\alpha_2\sqrt{I})$$
⁽¹⁷⁾

where the $\lambda_{MX}^{(n)}$ (*n* = 0, 1, 2) are the model coefficient parameters. The function *g*(*x*) is given by:

$$g(x) = \left(\frac{2}{x^2}\right) (1 - (1 + x)e^{-x})$$
(18)

In most cases, $\lambda_{MX}^{(2)}$ is set to zero and the parameter α_I is assigned a value of 2.0 (Pitzer, 1973). When *MX* comprises a 2:2 aqueous neutral electrolyte (and sometimes in other circumstances), $\lambda_{MX}^{(2)}$ is not set to zero, and α_2 and α_2 and are fixed at respective values of 1.4 and 12.0. Other values of α_2 and α_2 have occasionally been proposed (see Pitzer, 1987).

The second order interaction coefficient in the case of cation-cation, anion-anion, neutral-neutral, neutral-cation, and neutral-anion interactions is generally taken to be a constant. Thus, for such combinations one may write:

$$\lambda_{ij} = \lambda_{ij}^{(0)} \tag{19}$$

Pitzer (1975) modified his original treatment by adding a theoretical description for higher order electrostatic interactions. In the context of primitive interaction coefficients, this took the form of redefining the following second order interaction coefficients for the cation-cation and anion-anion combinations as:

$$\lambda_{MM'}(I) = {}^{S} \lambda_{MM'} + {}^{E} \lambda_{MM'}(I)$$
⁽²⁰⁾

$$\lambda_{XX'}(I) = {}^{S}\lambda_{XX'} + {}^{E}\lambda_{XX'}(I)$$
⁽²¹⁾

Here *M* and *M'* denote two cations and *X* and *X'* two anions. The description of the electrostatic term (${}^{E}\lambda_{MM'}(I)$ and ${}^{E}\lambda_{XX'}(I)$) are obtained from entirely from theoretical expressions; see Pitzer, 1975, and Chapter 3 of Wolery, 1992b). The original term (the "short range" term) can be written analogously to eq (19):

$${}^{S}\lambda_{MM'} = \lambda_{MM'}^{(0)}$$
(22)

$${}^{S}\lambda_{XX'} = \lambda_{XX'}^{(0)} \tag{23}$$

The temperature dependence of Pitzer interaction coefficients and interaction coefficient parameters (excluding *b*, α_1 , and α_2 , which are treated as constants) can be represented up to no more than 100°C by a Taylor's series truncated at first order, using the values of the coefficients and their first temperature derivatives at 25°C (see Silvester and Pitzer, 1978;Pitzer, 1978, 1987; Chapter 3 of Wolery, 1992b). Such a truncated Taylor's series has the general form:

$$x(T) = x_0 + \left(\frac{dx}{dT}\right)_{T_0}(T - T_0)$$
(24)

where x_0 is the value of the parameter at temperature T_0 (here 25°C). To obtain a better representation, one may use a Taylor's series truncated at second order. This has the general form:

$$x(T) = x_0 + \left(\frac{dx}{dT}\right)_{T_0}(T - T_0) + \frac{1}{2!}\left(\frac{d^2x}{dT^2}\right)_{T_0}(T - T_0)^2$$
(25)

This is presently the only means built into EQ3/6 for treating the temperature dependence of interaction coefficients and interaction coefficient parameters. Some other proposed methods but not yet implemented in EQ3/6 are discussed later in this chapter.

The use of the Pitzer's equations option requires a supporting data file consistent with these equations (e.g., **hmw** or **pit**). The model parameters required by EQ3/6 are:

- The A_{ϕ} Debye-Hückel parameter.
- 25°C values of the second order interaction coefficient parameters $\lambda_{ij}^{(0)}$, $\lambda_{ij}^{(1)}$, and $\lambda_{ij}^{(2)}$, the corresponding α_I and α_2 parameters for each *ij* pair, and the third order interaction coefficients μ_{ijk} . The parameter $\lambda_{ij}^{(2)}$ (and consequently α_2) is only used in certain cases; the parameter $\lambda_{ij}^{(1)}$ (and consequently α_I) is used in many, but not all cases.
- The first and second temperature derivatives of the interaction coefficient parameters λ⁽⁰⁾_{ij}, λ⁽¹⁾_{ij}, and λ⁽²⁾_{ij}, and the third order interaction coefficients μ_{ijk} (if calculations are to be made for temperatures other than 25°C). Derivatives are not required for λ⁽¹⁾_{ij}, and λ⁽²⁾_{ij} if these parameters themselves are not used for a given combination of solute species.

The A_{ϕ} parameter is represented in the usual grid format and is transformed into a set of interpolating polynomials by EQPT in the manner previously described for log *K* values and other kinds of Debye-Hückel parameters.

There are more of the $\lambda_{MX}^{(n)}$ and μ_{ijk} parameters than can be physically observed (Pitzer, 1973; see also Chapter 3 of Wolery, 1992b). These parameters can only be observed in certain combinations, which depend on the electrical charge types of the species. These combinations are equivalent to the observable interaction coefficients. There is a multiplicity of such coefficients, and the equation for the activity coefficient in complex mixtures becomes accordingly more complex (see Pitzer, 1979, 1987; Harvie, Møller, and Weare, 1984; and Felmy and Weare, 1986). However, relatively simple forms can be obtained for simple solutions, such as for a pure aqueous electrolyte (Pitzer, 1973, 1979, 1987), a mixture of two electrolytes with a common ion (Pitzer, 1973, 1975, 1979, 1987), and a mixture of an electrolyte and a neutral solute species (Pitzer, 1987). This facilitates much of the fitting of observed coefficients, and the data are consequently commonly reported in this form.

We consider first the parameters associated only with cations and anions. The parameters $\beta_{MX}^{(0)}$, $\beta_{MX}^{(1)}$, $\beta_{MX}^{(2)}$, and C_{MX}^{ϕ} , are observable in a pure solution of the aqueous neutral electrolyte comprised of cation *M* and anion *X*. The parameters ${}^{S}\theta_{MM'}$ and $\psi_{MM'X}$ are observable in a mixture of two aqueous neutral electrolytes, one comprised of cation *M* and anion *X*, the other of cation *M* and anion *X*. Similarly, the parameters ${}^{S}\theta_{XX'}$ and $\psi_{MXX'}$ are observable in a mixture of two aqueous neutral electrolytes, one comprised of cation *M* and anion *X*, the other of two aqueous neutral electrolytes, one comprised of cation *M* and anion *X*, the other of two aqueous neutral electrolytes, one comprised of cation *M* and anion *X*, the other of two aqueous neutral electrolytes, one comprised of cation *M* and anion *X*, the other of two aqueous neutral electrolytes, one comprised of cation *M* and anion *X*, the other of two aqueous neutral electrolytes, one comprised of cation *M* and anion *X*, the other of two aqueous neutral electrolytes, one comprised of the aqueous neutral electrolytes.

The relationship of these observable Pitzer parameters to the corresponding primitive Pitzer parameters is discussed in detail in Chapter 3 of Wolery (1992b). The observable parameters may be mapped to an equivalent set of primitive parameters by means of certain mapping relations. Those used by EQPT are the following:

$$\lambda_{MX}^{(n)} = \beta_{MX}^{(n)}$$
 for $n = 0, 2$ (26)

$$\lambda_{MM}^{(n)} = 0 \quad \text{for } n = 0, 2$$
 (27)

$$\lambda_{XX}^{(n)} = 0$$
 for $n = 0, 2$ (28)

$$\mu_{MMX} = \frac{1}{6} \left| \frac{z_M}{z_X} \right|^{\frac{1}{2}} C_{MX}^{\phi} \tag{29}$$

$$\mu_{MXX} = \frac{1}{6} \left| \frac{z_X}{z_M} \right|^{\frac{1}{2}} C_{MX}^{\phi} \tag{30}$$

$$\lambda_{MM'}^{(0)} = {}^{S} \Theta_{MM'} \tag{31}$$

$$\lambda_{XX'}^{(0)} = {}^{S} \Theta_{XX'} \tag{32}$$

$$\mu_{MM'X} = \frac{1}{6} \left(\Psi_{MM'X} + \left(\frac{3z_M}{z_M} \right) \mu_{MMX} + \left(\frac{3z_M}{z_{M'}} \right) \mu_{M'M'X} \right)$$
(33)

$$\mu_{MXX'} = \frac{1}{6} \left(\Psi_{MXX'} + \left(\frac{3z_{X'}}{z_X} \right) \mu_{MXX} + \left(\frac{3z_X}{z_{X'}} \right) \mu_{MX'X'} \right)$$
(34)

The temperature derivatives of the observable Pitzer parameters may be mapped to the temperature derivatives of the corresponding conventional primitive parameters using the temperature derivatives of the mapping relations for the parameters themselves. For example, differentiation of eq (26) gives:

$$\frac{d\lambda_{MX}^{(n)}}{dT} = \frac{d\beta_{MX}^{(n)}}{dT} \quad \text{for } n = 0, 2$$
(35)

$$\frac{d^2 \lambda_{MX}^{(n)}}{dT^2} = \frac{d^2 \beta_{MX}^{(n)}}{dT^2} \quad \text{for } n = 0, 2 \tag{36}$$

The mapping relations for the second derivatives are analogous.

Harvie, Møller, and Weare (1984) used Pitzer's equations to construct a model of all of the major components of seawater at 25°C. They modified the equations for electrolyte systems to include some provision for neutral species-ion interactions. Felmy and Weare (1986) took the equations further in this direction when they extended the Harvie, Møller, and Weare model to include borate as a component. These investigators introduced the following new observable parameters: λ_{NM} and λ_{NX} , and ζ_{NMX} . Here N denotes a neutral species. The terms in λ_{NM} and λ_{NX} were introduced by Harvie, Møller, and Weare (1984) in order to treat the species $CO_{2(aq)}$ in their model of the "sea salt" system. They are treated as constants. To deal with the fact that they are only observable in combination, Harvie, Møller, and Weare (1984) adopted the following convention:

$$\lambda_{N,H^+} = 0 \tag{37}$$

We note that Clegg and Brimblecombe (1989, 1990) use a different convention:

$$\lambda_{N, Cl} = 0 \tag{38}$$

It is important to follow a single convention in any data file. At the present time, that of Harvie, Møller, and Weare (1984) is used in the **hmw** data file. The present version of the **pit** data file contains no λ_{NM} or λ_{NX} parameters.

The ζ_{NMX} parameter is an observable third order coefficient. It was developed by Felmy and Weare (1986) in order to account for interactions involving the species $B(OH)_{3(aq)}$. This parameter can be mapped into primitive equivalents by means of the following mapping conventions (see Chapter 3 of Wolery, 1992b):

$$\mu_{NMX} = \frac{\zeta_{NMX}}{6} \tag{39}$$

$$\mu_{NMM} = 0 \tag{40}$$

$$\mu_{NXX} = 0 \tag{41}$$

There are additional possible Pitzer coefficients for solutions containing neutral solute species (see Chapter 3 of Wolery, 1992b). Among these, the most likely to be needed are the second order interaction coefficients λ_{NN} and $\lambda_{NN'}$ (here N' denotes a second neutral species). These coefficients are directly observable.

The temperature dependence of observable and conventional primitive coefficients involving neutral solute species can be handled analogously to that manner previously described for coefficients and coefficient parameters involving only cations and anions; e.g., using tabulated first and second derivatives. The mapping functions for the derivatives are again analogous to those for the coefficients themselves.

The relevant EQ3/6 data files contain two superblocks of observable Pitzer coefficient data (a superblock is a collection of similar data blocks). The first superblock consists of blocks containing the data observable in solutions of pure aqueous neutral electrolytes: $\beta_{MX}^{(0)}$, $\beta_{MX}^{(1)}$, $\beta_{MX}^{(2)}$, and

 C_{MX}^{ϕ} . Each of these blocks also contains the corresponding values of α_1 and α_2 . The second superblock consists of blocks containing the data observable in mixtures of two aqueous neutral electrolytes containing a common ion: ${}^{S}\theta_{MM'}$ and $\psi_{MM'X}$, or ${}^{S}\theta_{XX'}$ and $\psi_{MXX'}$. The data in either type of superblock in represented by the 25°C value of the relevant parameters, plus the first and second temperature derivatives of these parameters at the same temperature (note: α_1 and α_2 are taken to be independent of temperature).

We note a problem concerning ${}^{S}\theta_{MM'}$ and its temperature derivatives. A potentially different value can be obtained by fitting measurements on more than one mixture of two aqueous electrolytes containing the two cations *M* and *M'*. The same problem holds for ${}^{S}\theta_{XX'}$ and its temperature derivatives. This is particularly a problem in that potentially different values of the theta coefficients and their temperature derivatives may appear on the data file for each relevant mixture. EQPT deals with this by taking the average of such parameters. Strictly speaking, the data file should not contain the different values for the any given theta coefficient. The same rule applies to its first and second order temperature derivatives.

The relevant EQ3/6 data files also contain a flag string which alerts EQ3NR or EQ6 to use or ignore the higher order electrostatic model proposed by Pitzer (1975). This is also written on the **data1** file produced by EQPT. Note that the fitted values of the theta and psi parameters involving two cations not of the same charge or two anions not of the same charge according to whether or not this model is included.

The present version of EQPT was actually designed to handle parameters involving only cations and anions. However, it can deal with λ_{NM} by composing a pure aqueous neutral electrolyte block for a fictive electrolyte. Here *N* and *M* are specified as the pair of "ions" composing the "electrolyte." The value of λ_{NM} is entered in the field for $\beta_{NM}^{(0)}$. The remaining parameter fields in this block should be left blank or filled with zeros. The λ_{NX} parameter can be treated likewise. Coefficients of the type λ_{NN} and $\lambda_{NN'}$ can be dealt with by using the same trick used to deal with λ_{NM} . In this case, the fictive electrolyte consists of two neutral species (*N* and *N*, or *N* and *N'*). However, no examples of such coefficients appear on the **hmw** data file or the existing **pit** data file. The present version of EQPT can not deal with the ζ_{NMX} parameter, however. The two superblocks for observable Pitzer parameters and their temperature derivatives are mapped by EQPT to corresponding superblocks on the **data1** file containing the corresponding conventional primitive parameters and their temperature derivatives. EQ3NR and EQ6 then use these data to calculate values of the relevant conventional primitive parameters at the desired temperature. Note that if the derivatives are missing, no temperature correction is made.

The **hmw** data file has nominal lower and upper temperature limits which are both set to 25°C. It contains no temperature derivative data. It should not be used to make calculations at temperature other than 25°C. EQ3NR and EQ6 will write warnings if the nominal temperature limits are exceeded. This data file also employs the higher order electrostatic model proposed by Pitzer (1975).

The **pit** data file has nominal lower and upper temperature limits of 0°C and 100°C, respectively. It contains a fair amount of data for first temperature derivatives. This data file does not employ the higher order electrostatic model proposed by Pitzer (1975).

The temperature dependency has been expressed in various more recent studies by a variety of different temperature functions, most of which require only 5-7 parameters per observable interaction coefficient. Pabalan and Pitzer (1987) used such equations to develop a model for the system $Na-K-Mg-Cl-SO_4-OH-H_2O$ which appears to be generally valid up to about 200°C. Pabalan and Pitzer (1988) used equations of this type built a model for the system $Na-Cl-SO_4-OH-H_2O$ that extends to 300°C. Greenberg and Møller (1989), using an elaborate compound temperature function, have constructed a model for the $Na-K-Ca-Cl-SO_4-H_2O$ system that is valid from 0-250°C. More recently, Spencer, Møller, and Weare (1990) have used a more compact equation to develop a model for the $Na-K-Ca-Mg-Cl-SO_4-H_2O$ system at temperatures in the range -60-25°C. The present version of EQPT has no capability to deal with these kinds of representations of the temperature dependence.

3.2.2. Activity Coefficients of Solid Solution Components

In general, the activity coefficient of a solid solution component depends on the composition of the solid solution. This in turn is normally expressed in terms of the mole fractions of the endmember components (for example, calcite $[CaCO_3]$ and magnesite $[MgCO_3]$ in magnesian calcite $[(Ca,Mg)CO_3]$). There are two categories of activity coefficient models, molecular-mixing models and site-mixing models (cf. Chapter 4 of the EQ3NR Theoretical Manual and User's Guide, Wolery, 1992b).

In molecular-mixing models, the activity coefficient of an end-member component is unity in the ideal case. In the non-ideal case, one generally utilizes a model which describes the activity coefficient via a set of interaction coefficients similar to those employed in Pitzer's equations for the activity coefficients of aqueous species.

In site-mixing models, explicit account is taken of the fact that mixing of ions occurs on welldefined sites in the crystal structure (see for example Wood and Fraser, 1977, or Nordstrom and Munoz, 1985). Vacancies may be present on a site, and be created or destroyed by substitutions of one ion for another of different electrical charge. In site-mixing models, the concept of ideality is modified from that appropriate to molecular mixing, though still based on the mole fractions of components. It is possible to utilize as the components species such as ions, vacancies, and framework moieties instead of end members, but the more common practice is to continue using end-member components. This is followed in the present version of EQ3/6. However, the activity coefficient of an end-member component in an ideal site mixing model may have a value other than unity. A site-mixing model will appear to be ideal in this sense only if there is only one site, an ion substitutes for others of the same charge type, and vacancies are not present on the site. Site mixing then effectively reduces to molecular mixing.

Nearly all of the existing site-mixing models of solid solutions are ideal in the site-mixing sense (see for example Viani and Bruton, 1992). The only parameters of such models are site-mixing parameters. It is possible to consider site-mixing models which are non-ideal even in the site-mixing sense. These would be described by both site-mixing parameters and interaction coefficients. No models of this type are presently treated in EQ3/6.

In EQ3/6, all solid solution models are defined on the supporting data file (see Chapter 4). At present (through the R16 set of data 2files), only the **com** file contains any solid solutions. Most of these are treated with a simple ideal site-mixing model (the only exception concerns olivine, which is treated according to a regular solution model). The actual types of models used on the data file are defined by the **jsol** flag array. The corresponding parameters (site-mixing parameters, interaction coefficients, and parameters used to compute interaction coefficients) are stored on the data file in the **apx** array. The elements of this array are represented below as p_{kw} .

The various models presently treated in EQ3/6 are briefly discussed in the following sections. To avoid confusion, we will often write the activity, mole fraction, activity coefficient, and related parameters of a solid solution component with a " $\sigma\psi$ " subscript in place of "*i*." Here σ will denote the component itself (takes the place of "*i*"), and ψ the solid solution (in order to be completely explicit about which solid solution is being addressed).

3.2.2.1. Ideal Solution, with One Optional Site-Mixing Parameter

The first activity coefficient model for solid solutions in EQ3/6 is for an ideal solution in either the molecular-mixing sense or a limited site-mixing model in which mixing is confined to one site and vacancies are ignored. The former is a special case of the latter. This model corresponds to **jsol** = 1 and is characterized by the equation (Wood and Fraser, 1977; Viani and Bruton, 1992):

$$a_{\sigma\psi} = x_{\sigma\psi}^{N_{\psi}} \tag{42}$$

where N_{Ψ} is the site mixing parameter. This formulation is equivalent to:

$$log\lambda_{\sigma\Psi} = (N_{\Psi} - 1)logx_{\sigma\Psi}$$
(43)

If $N_{\psi} = 1$, the above model is mathematically equivalent to an ideal molecular-mixing model $(\log \lambda_{\sigma \psi} = 0)$.

The N_{ψ} parameter is stoichiometric in nature. In essence, it is the number of formula units of the site on which mixing occurs per formula unit of the solid solution framework. In principle, the

formula for all the end-member components of a solid solution can be written so as to yield N_{ψ}

= 1, hence
$$log\lambda_{\sigma\Psi} = 0$$
.

In the case of heterovalent single-site solid solutions such as clays and zeolites, vacancies are involved. In order to simplify the solution model and preserve the simple relationship defined by eqs (42) and (43), Viani and Bruton (1992) have chosen to treat such solid solutions according to a model in which the mixing entities are ions or ion-vacancy complexes. Thus, two sodium ion entities might mix with a calcium ion-vacancy entity.

The N_{Ψ} parameter is obtained from the parameters read from the data file according to:

$$N_{\Psi} = p_{7\Psi} \tag{44}$$

3.2.2.2. Third-Order Maclaurin Model for a Binary Solution

The third-order Maclaurin model for a binary solution corresponds to $\mathbf{jsol} = 2$. It is taken from Helgeson et al. (1970). The activity coefficients of the two end-member components are given by:

$$log\lambda_{1\psi} = \frac{1}{2.303RT} \left[-\left(\frac{W_{2\psi}}{2}\right) x_2^2 - \left(\frac{W_{3\psi}}{3}\right) x_2^3 \right]$$
(45)

$$\log \lambda_{2\psi} = \frac{1}{2.303RT} \left[-\left(\frac{W_{2\psi} + W_{3\psi}}{2}\right) x_1^2 + \left(\frac{W_{3\psi}}{3}\right) x_1^3 + \left(W_{1\psi} + \frac{W_{2\psi}}{2} + \frac{W_{3\psi}}{6}\right) \right]$$
(46)

Here $W_{1\psi}$, $W_{2\psi}$, and $W_{3\psi}$ are interaction coefficients. There are no site-mixing parameters.

The formulation represented by eqs (45) and (46) is highly unsymmetrical. In order to satisfy the condition that $log \lambda_{2\psi} \rightarrow 0$ as $x_1 \rightarrow 0$, the interaction coefficients are required to satisfy the relation:

$$W_{1\psi} = -\frac{W_{2\psi}}{2} - \frac{W_{3\psi}}{6} \tag{47}$$

The interaction coefficients are obtained from the parameters read from the data file according to:

$$W_{1\psi} = p_{1\psi} \tag{48}$$

$$W_{2\psi} = p_{2\psi} \tag{49}$$

$$W_{3\psi} = p_{3\psi} \tag{50}$$

However, $W_{I\psi}$ is actually recalculated using eq (47).

3.2.2.3. Regular Solution Model for a Binary Solution

The regular solution model for a binary solution corresponds to $\mathbf{jsol} = 3$. It is also called a parabolic Maclaurin model. For a discussion of this model, see Saxena (1973, p. 11-12). The activity coefficients of the two end-member components are given by:

$$\log \lambda_{1\psi} = \frac{1}{2.303RT} W_{\psi} x_2^2$$
(51)

$$\log \lambda_{2\psi} = \frac{1}{2.303RT} W_{\psi} x_1^2$$
 (52)

Here W_{ψ} is the single interaction coefficient. There are no site-mixing parameters. This formulation is symmetrical.

The interaction coefficients are obtained from the parameters read from the data file according to:

$$W_{\psi} = p_{1\psi} + p_{2\psi}T + p_{3\psi}P \tag{53}$$

Thus, the interaction coefficient in this model can be treated as a function of temperature and pressure. On the **com** data file in the R10 and R16 sets, there is a regular solution model for the solid solution olivine. The $p_{2\psi}$ and $p_{3\psi}$ parameters are set to zero, so the interaction coefficient is actually treated as a constant. A non-unit site-mixing parameter is also given in the $p_{7\psi}$ parameter, but this is not used.

3.2.2.4. Cubic Maclaurin Model for a Binary Solution

The cubic Maclaurin model for a binary solution corresponds to jsol = 4. For a discussion of this model, see Saxena (1973, p. 16). The activity coefficients of the two end-member components are given by:

$$log\lambda_{1\psi} = \frac{1}{2.303RT} [(2W_{2\psi} - W_{1\psi})x_2^2 + 2(W_{1\psi} - W_{2\psi})x_2^3]$$
(54)

$$\log \lambda_{2\psi} = \frac{1}{2.303RT} [(2W_{1\psi} - W_{2\psi})x_1^2 + 2(W_{2\psi} - W_{1\psi})x_1^3]$$
(55)

Here $W_{I\psi}$ and $W_{2\psi}$ are interaction coefficients. There are no site-mixing parameters. This formulation is asymmetrical.

The interaction coefficients are obtained from the parameters read from the data file according to:

$$W_{1\psi} = p_{1\psi} + p_{2\psi}T + p_{3\psi}P$$
(56)

$$W_{2\psi} = p_{4\psi} + p_{5\psi}T + p_{6\psi}P$$
(57)

3.2.2.5. Guggenheim Polynomial Model for a Binary Solution

The Guggenheim polynomial model for a binary solution corresponds to $\mathbf{jsol} = 5$. For a discussion of this model, see Saxena (1973, p. 14-15). The activity coefficients of the two end-member components are given by:

$$log\lambda_{1\psi} = \frac{1}{2.303RT} [(W_{1\psi} + 3W_{2\psi} + 5W_{3\psi})x_2^2 + (-4W_{2\psi} - 16W_{3\psi})x_2^3 + 12W_{3\psi}x_2^4]$$
(58)

$$log\lambda_{2\psi} = \frac{1}{2.303RT} [(W_{1\psi} - 3W_{2\psi} + 5W_{3\psi})x_1^2 + (4W_{2\psi} - 16W_{3\psi})x_1^3 + 12W_{3\psi}x_1^4]$$
(59)

Here $W_{I\psi}$, $W_{2\psi}$, and $W_{3\psi}$ are interaction coefficients. There are no site-mixing parameters. This formulation is asymmetrical.

The interaction coefficients are obtained from the parameters read from the data file according to:

$$W_{1\psi} = p_{1\psi} + p_{2\psi}T + p_{3\psi}T^2$$
(60)

$$W_{2\psi} = p_{4\psi} + p_{5\psi}T + p_{6\psi}T^2$$
(61)

$$W_{3\psi} = p_{7\psi} + p_{8\psi}T + p_{9\psi}T^2$$
(62)

The full form of this model can be used in the present version of EQ3/6, although the parameters p_{kw} for $k \ge 7$ are now intended to be reserved for site-mixing parameters.

3.2.2.6. Regular Solution Model for a Ternary Solution

The regular solution model for a ternary solution corresponds to $\mathbf{jsol} = 6$. For a discussion of this model, see Prigogine and Defay (1954, p. 257). The activity coefficients of the three end-member components are given by:

$$\log \lambda_{1\psi} = \frac{1}{2.303RT} [W_{12\psi} x_2^2 + W_{13\psi} x_3^2 + (W_{12\psi} - W_{23\psi} + W_{13\psi}) x_2 x_3]$$
(63)

$$log\lambda_{2\psi} = \frac{1}{2.303RT} [W_{12\psi}x_1^2 + W_{23\psi}x_3^2 + (W_{12\psi} - W_{13\psi} + W_{23\psi})x_1x_3]$$
(64)

$$\log \lambda_{3\psi} = \frac{1}{2.303RT} [W_{13\psi} x_1^2 + W_{23\psi} x_2^2 + (W_{13\psi} - W_{12\psi} + W_{23\psi}) x_1 x_2]$$
(65)

Here $W_{I\psi}$, $W_{2\psi}$, and $W_{3\psi}$ are interaction coefficients. There are no site-mixing parameters. This formulation is symmetrical.

The interaction coefficients are obtained from the parameters read from the data file according to:

$$W_{12\psi} = p_{1\psi} \tag{66}$$

$$W_{13\psi} = p_{2\psi} \tag{67}$$

$$W_{23\psi} = p_{3\psi} \tag{68}$$

4. Data File Contents and Structures

The **data0** data files are represented by two archetypes. The first, called the **com** archetype is used for data files intended to support calculations using simple extended Debye-Hückel models for the activity coefficients of the aqueous species. This archetype pertains to the **com**, **sup**, and **nea** data files. The **hmw** archetype is used for data files intended to support calculations using Pitzer's (1973, 1975, 1979, 1987) equations. It pertains to the **hmw** and **pit** data files. The two archetypes differ in the types of parameters for computing activity coefficients of aqueous species. Otherwise, they are essentially the same.

Archetypes for the **data1** file structures closely parallel those for the corresponding **data0** files. The **data1** file is an unformatted file, so no formats (in the usual sense) are involved, and the file can not be printed or displayed in any meaningful fashion. To assist debugging, EQPT writes a corresponding data file called **data1f**, which is a formatted equivalent of **data1**.

The contents of **data1** are identical to those of **data0**, with the following exceptions. Data given on a standard temperature grid are replaced by the coefficients of interpolating polynomials which EQPT fits to such data grids. Also, in the case of data files of the **hmw** archetype, observable Pitzer coefficient parameters are mapped to an equivalent set of conventional primitive parameters. The temperature derivatives of these observable parameters are mapped similarly to those of the conventional primitive equivalents. All of these data transformations were discussed in detail in Chapter 3.

4.1. The com Archetype for the data0 File

The basic structure of the **com** archetype for **data0** file is given in Figure 2. The data file begins with a one-line header of the form:

```
data0.com.R10
```

The first five letters on this line must be **data0**. If they are not EQPT will terminate with an error message. Letters 7-9 must contain the three letter data file key string (**com**, **nea**, **sup**, **hmw**, or **pit**). The stage number (**R10** in the above example) is optional as far as EQPT is concerned. However, it should appear to assist in configuration management, as it will be written on the **data1** file by EQPT, and thence by EQ3NR and EQ6 on their **output** files.

The header is followed by a title, which may consist of up to 70 lines (see **ntitpa** in Appendix A) of descriptive text. The title on the **data0.com.R10** data file is:

The "+-----" in the final line in this example is a block terminator used throughout the data file. EQPT writes the data file title on the **data1** file. EQ3NR and EQ6 read this title and write it on their **output** files. Any changes to the data file made by users may be noted or cited in this title. Users outside the LLNL Data Base Development Task who make any such changes are requested to make some kind of notation on modified data files so as to distinguish them from those supplied by LLNL.

eader (data0, data file key, stage number
Title
"data0" parameters block:
• Temperature limits (°C)
• Temperature (°C), standard grid
• Pressure (bars), standard grid
• $A_{\gamma, 10}$, standard grid
• B_{γ} , standard grid
• <i>B</i> , standard grid
• Drummond (1981) <i>C</i> , <i>F</i> , <i>G</i> , <i>E</i> , and <i>H</i>
coefficients
• $log K_{Eh}$, standard grid
"bdot" parameters block:
-
• Aqueous species name, a_i , insgfl flag
Chemical elements block:
• Elements, atomic weights
Aqueous species superblock:
• Strict basis species
• Auxiliary basis species, with <i>log K</i>
on the standard grid
-
• Non-basis species, with <i>log K</i>
on the standard grid
Pure minerals superblock:
• Pure minerals, with <i>log K</i>
on the standard grid
Pure liquids superblock:
• Pure liquids, with log K
on the standard grid
Gas species superblock:
• Gas species, with log K on the standard
grid
gnu
Solid solutions superblock:
Solid solutions
References block:
References

Figure 2. The basic structure of the data0 file for the com archetype.

The title is followed by a block of data for "data0" parameters. These include the nominal temperature limits (°C) for the application of the data file, the temperatures (°C) on the standard grid, the pressure (bars) on the standard temperature grid, the $A_{\gamma, I0}$ and B_{γ} Debye-Hückel parameters

on the standard temperature grid, the extended Debye-Hückel \dot{B} on the standard temperature grid, the five Drummond 1981 coefficients (*C*, *F*, *G*, *E*, and *H*) needed to compute the activity coefficient of aqueous CO_2 (see Chapter 3 of Wolery, 1992b), and $log K_{Eh}$ on the standard temperature grid. The parameter $log K_{Eh}$ is the equilibrium constant of the special reaction (9) used in EQ3NR and EQ6 to relate seondary redox variables (*Eh*, *pe*) from the primary redox variable, teh oxygen fugacity. This reaction itself is not written on the data file. It is hard-wired into EQ3NR and EQ6.

data0 parameters _____ temperature limits (a) 0.0000 300.0000 (5x,2f10.4) temperatures 0.0100 25.0000 60.0000 100.0000 (5x,4f10.4) 150.0000 200.0000 250.0000 300.0000 pressures 1.0132 1.0132 1.0132 1.0132 4.7572 15.5365 39.7365 85.8378 debye huckel a (adh) 0.4939 0.5114 0.5465 0.5995 0.6855 0.7994 0.9593 1.2180 debye huckel b (bdh) 0.3288 0.3253 0.3346 0.3421 0.3525 0.3766 0.3925 0.3639 bdot 0.01 0.0470 for t 0.0174 0.0440 0.0460 0.0340 0.0470 0.0000 (coefficients for the Drummond (1981) polynomial) cco2 -1.0312 0.0012806 (5x,f10.4,11x,f12.7) 255.9 (10x.f5.1,11x,f12.4) 0.4445 -0.001606 (5x,f10.6) log k for eh reaction (5x,4f10.4)

The "data0" parameters block in data0.com.R10 is as follows:

The "data0 parameters" on the first line is a block header flag. EQPT actually uses the "temperature limits" string to position the data file for reading this block. The other individual header strings are not read. The formats for reading the data are superimposed to the right in the above example where they appear in bold italic. These format strings do not appear in the data file itself. Formats which obviously repeat are not marked. We will continue this practice of giving formats in this manner in the examples to follow. Note the usage here of the standard pressure grid. The pressure of "4.7572" bars is the pressure at "150.0000" degrees (C).

This is followed by the "bdot" parameters block. This block consists of its own header and a list of the hard core diameter (a_i) and **insgfl** flag switch. This block in **data0.com.R10** is as follows:

bdot parameters					(a)
<pre>* species name (uo2)3(co3)6(6-) np(co3)5(6-) u(co3)5(6-)</pre>	azer0 4.0000 4.0000 4.0000	neutral 0 0 0	ion	type	(a24,7x,f7.1,4x,i2)
(Material Deleted)					
th6(oh)15(9+) +	6.0000	0			

Here "azer0" is the hard core diameter, and the "neutral ion type" is the **insgfl** flag. The former is given in units of cubic angstroms. The latter is meaningful only in the case of neutral species. If **insgfl** = -1, the activity coefficient of the species is taken to be the value for aqueous CO_2 in pure aqueous NaCl, as computed from eq (12) (see Chapter 3). This is appropriate for nonpolar species, such as most of the dissolved gas species. If **insgfl** = 0, the activity coefficient is set to unity. This is more appropriate for polar species, such as the $MgSO_{4(aq)}$ ion pair. The name of an aqueous species appearing in this block must be identical to its name in the corresponding species block, which appears below this block. EQPT writes the data in this block asis onto the **data1** file. The line beginning with "* species name" is a comment line. Comment lines begin with an asterisk in column one. EQPT actually reads a copy of the **data0** file from which the comment lines have been stripped.

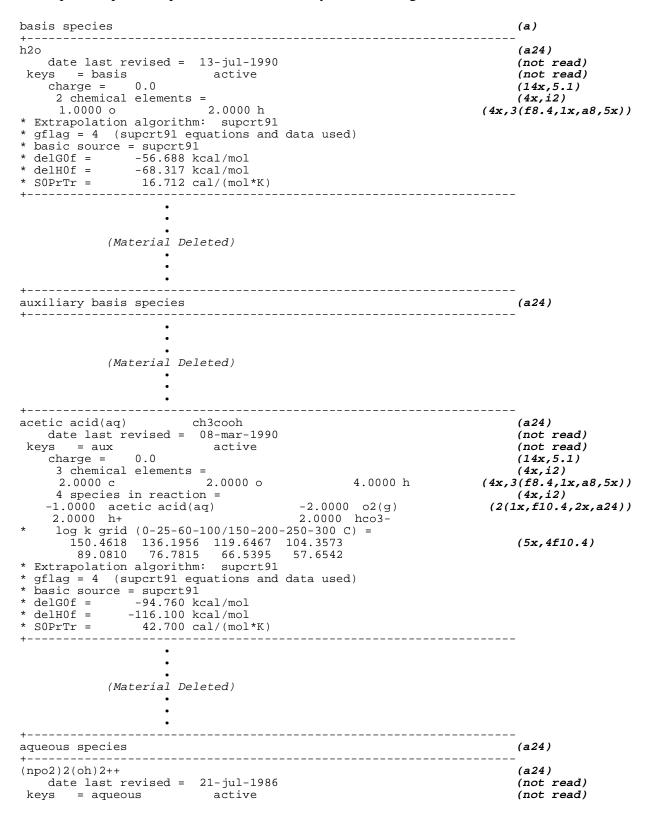
Next is the chemical elements block,. It consists of a block header followed by the names of the chemical elements (represented by the standard symbols) and their atomic weights (grams per mole). It is illustrated below by the block from **data0.com.R10**:

elements	(a)		
+ 0	15.99940		
ag	107.86820		(a8,f10.5)
al	26.98154		
	•		
	•		
	•		
	(Material	Deleted)	
	•		
	•		
	•		
zr	91.22400		
+			

This is followed by the aqueous species superblock. This is comprised of a data block for each aqueous species. The structure of this superblock is complicated somewhat in that these data blocks are organized into three "sub-superblocks," the first for strict basis species, the second for auxiliary basis species, and the third for non-basis aqueous species. Furthermore, water ("h2o") must be the first strict basis species. The fictive aqueous redox species "o2(g)" must be the last strict basis species. Each strict basis species except "o2(g)" must correspond to a chemical element in the chemical element solock. They should also appear in corresponding order. Note that water corresponds to the element oxygen. The strict basis species sub-superblock is terminated by a short block containing the string "auxiliary basis species" in place of a species

name. The auxiliary basis species sub-superblock is similarly terminated by a short block containing the string "aqueous species".

The aqueous species superblock is illustrated by the following, taken from data0.com.R10:



```
charge = 2.0
                                                                     (14x, 5.1)
    3 chemical elements =
                                                                     (4x,i2)
    3 Chemice.

2.0000 h

4 species in reaction =

1 0000 (npo2)2(oh)2++

2.0000 h+

2.0000 h+

2.0000 npo2++

2.0000 c) =
                                             6.0000 o
                                                               (4x, 3(f8.4, 1x, a8, 5x))
                                                                     (4x,i2)
   -1.0000 (npo2)2(oh)2++
2.0000 h2o
                                                                (2(1x,f10.4,2x,a24))
    log k grid (0-25-60-100/150-200-250-300 C) =
*
      500.0000 6.4000 5.6000 5.0000
4.6000 500.0000 500.0000 500.0000
                                                                     (5x,4f10.4)
* gflag = 3 (reported logk data used)
* logk source = 841em
   calculated g-h-s values:
* delGOf = -485.046 kcal/mol
* delHOf = -537.089 kcal/mol
* SOPrTr = -3.346 cal/(mol*K)
+------
                                  _____
            (Material Deleted)
```

Note that the species blocks illustrated here each contain a number of comment lines. Also, the blocks for non-basis aqueous species block and auxiliary basis species share the same format. Note that "500.0000" is used in the *log K* grid to mean "no data." This is illustrated in the data block for "(npo2)2(oh)2++".

The pure minerals superblock follows the aqueous species superblock. It is similar, but has no sub-superblocks and no special ordering restrictions. It is illustrated by the following, taken from **data0.com.R10**:

```
solids
                                                                            (a24)
                     _____
(pb(oh)2)3.pbcl2 (pb(oh)2)3.pblcl2
                                                                             (a24)
date last revised = 24-aug-1989
keys = solid active
VOPrTr = 0.000 cm**3/mol (source =
                                                                             (not read)
                                                                  (not read)
(16x,f9.3)
     4 chemical elements =
                                                                             (4x,i2)
                                                  (4x,i2)
6.0000 h (4x,3(f8.4,1x,a8,5x))
                            4.0000 pb
     2.0000 cl
   6.0000 o

5 species in data0 reaction

-1.0000 (pb(oh)2)3.pbcl2 -6.0000 h+

4.0000 pb++
      6.0000 o
                                                                             (4x, i2)
                                                                      (2(1x,f10.4,2x,a24))
     6.0000 h2o
     \log k \text{ grid } (0-25-60-100/150-200-250-300 \text{ C}) =

        500.0000
        17.2793
        500.0000
        500.0000

        500.0000
        500.0000
        500.0000
        500.0000

                                                                            (5x,4f10.4)
* gflag = 1 (reported delGOf used)
* basic source = 82wag/eva
* delGOf = -1682.600 kj/mol
* delHOf = 2092.000 kj/mol
* SOPrTr = 2092.000 j/(mol*K)
+------
                                  _____
             (Material Deleted)
                   _____
 _____
```

The pure liquids superblock follows. The format of a pure liquid block is identical to that in a pure mineral block. The pure liquids superblock is read by EQPT, but no corresponding data are written on the **data1** file. This is because EQ3NR and EQ6 have no capability for handling pure liquid species. Note that water does not appear in this superblock. This superblock is illustrated by the following, taken from **data0.com.R10**:

liquids _____ +---br2 date last revised = 18-may-1990 ys = liquid refstate active (see above for formats) = liquid refstate active VOPrTr = 0.000 cm**3/mol (source = keys) 1 chemical elements = 2.0000 br 5 species in data0 reaction -1.0000 h2o -1.0000 br2 0.5000 2.0000 o2(q) br-2.0000 h+ log k grid (0-25-60-100/150-200-250-300 C) = * -5.8592 -5.0927 -4.4059 -3.9728 -3.7893 -3.9054 -4.2985 -5.0485 * Extrapolation algorithm: cp integration * gflag = 1 (reported delGOf used) basic source = 89cox/wag * delGOf = 0.000 kj/mol * delHOf = 0.000 kj/mol * SOPTTr = 152.210 j/(mol*K) cp source = 79rob/hem units = jou T**0 0.36060000E+02 * cp source = 79rob/hem units = jou T**0 0.38426000E+02 T**-0.5 -0.22423000E+02 $\begin{array}{rcrcr} 1 & 1 & -0.5 & -0.22425000 \pm 102 \\ T^{**-2} & -0.95885000 \pm 105 \\ T^{**2} & 0.13663000 \pm -06 \\ Tlimit & = 1526.850 \ C \end{array}$ * * _____ (Material Deleted)

The gas species superblock follows. The format of a gas species block is identical to that in a pure mineral block. This superblock is illustrated by the following, taken from **data0.com.R10**:

```
gases
          _____
aq(q)
   date last revised = 05-apr-1988
                                                         (see above for formats)
     = gas active
VOPrTr = 0.000 cm**3/mol (source =
keys
                                                         )
    1 chemical elements =
     1.0000 ag
    5 species in data0 reaction
   -1.0000 ag(g)
                                     -1.0000 h+
   -0.2500
                                      0.5000 h2o
           o2(g)
    1.0000 ag+
    log k grid (0-25-60-100/150-200-250-300 C) =
*
       55.5477
                50.3736 44.4663 39.1150
                29.8254 500.0000 500.0000
       33.8983
* Extrapolation algorithm: constant enthalpy approximation
 gflag = 2 (calculated delGOf(delHOf,SOPrTr) used)
 basic source = 89cox/wag
* delGOf = 246.040 kj/mol
```

The solid solutions superblock follows the gas species superblock. A solid solution block consists of the name of the solid solution phase, a list of end-member component species, a mixing law to define the activity coefficients of these species, and a set of parameters for this mixing law. A set of site parameters may also be specified. Each end-member component must be represented on the data file as a pure mineral. For each end member, there is also specified an upper limit to its mole fraction in the solid solution. This superblock is illustrated by the following, taken from **data0.com.R10**:

solid solutions (a24) _____ (Material Deleted) _____ carbonate-calcite (ca,mn,zn,mg,fe,sr)co3 (a24) date last revised = 22-dec-1987 (not read) ideal active (not read) keys = ss 6 end members (i3)

 end members
 (i3)

 1.0000 calcite
 1.0000 magnesite
 (5x,2(f6.3,2x,a24,5x))

 1.0000 rhodochrosite
 1.0000 siderite
 (5x,2(f6.3,2x,a24,5x))

 1.0000 smithsonite
 1.0000 strontianite
 (5x,2(f6.3,2x,a24,5x))

 type = 1 (10x,i1) 0 model parameters (i3) 1 site parameters 1.000 0.000 0.000 0.000 0.000 0.000 (i3) (6£6.3) (Material Deleted) _____

Of the present five data files, only the **com** file contains any solid solutions. These are all ideal ("type = 1"), except for olivine, which is treated as a binary regular solution ("type = 3"). The "model parameters" are the set of interaction coefficient parameters. For an ideal solution, there are none. If there were any such parameters, they would be entered below "model parameters" using the format illustrated for "site parameters". Both kinds of parameters are currently stored in different parts of the same array (**apx**, see Chapter 3). In current practice, one site parameter is declared for each solid solution, regardless of type. This is stored as **apx**(7,**nx**) for the **nx**-th solid solution. If the solution is not ideal, however, the number given as a site parameter is not used as such (ion most cases is simply not used). It and other members in

the site parameter range of the **apx** array may be used instead to store additional interaction coefficient parameters.

The references block is the last part of the data file. It contains references. It is terminated by a line beginning with "stop.". It is illustrated by the following, taken from **data0.com.R10**:

EQPT does not write the contents of this block onto the **data1** file. This block is present only to provide a means of documenting the data present on the data file.

4.2. The hmw Archetype for the data0 File

The basic structure of the **hmw** archetype is illustrated in Figure 3 (compare with Figure 2). This is identical to the **com** archetype, except for two differences. The "data0" parameters block has a slightly different content, and the "bdot" parameters block is replaced by two superblocks, one for Pitzer coefficient data pertaining to pure aqueous neutral electrolytes, the other to such data pertaining to mixtures of two aqueous neutral electrolytes containing a common ion.

The "data0" parameters block in data0.hmw.R10 is as follows:

Note that A_{ϕ} appears in place of $A_{\gamma, 10}$. B_{γ} , \dot{B} , and the *C*, *F*, *G*, *E*, and *H* coefficients from Drummond (1981) do not appear in this block in this archetype. Although A_{ϕ} and $A_{\gamma, 10}$ are related by theory, the former should not be calculated from the latter, as slight but significant differences in the value of Debye-Hückel parameters occur from model to model (see Chapter 3), hence also from data file to data file. This is also why the values of the Debye-Hückel parameters are written on the data files instead of being hard-wired into EQ3NR and EQ6.

Header (data0 , data file key, stage number))
Title	
"data0" parameters block:	
• Temperature limits (°C)	
• Temperature (°C), standard grid	
• Pressure (bars), standard grid	
• A_{ϕ} , standard grid	
• $log K_{Eh}$, standard grid	
Pure electrolyte parameters superblock: • Species pairs; $\beta_{MX}^{(n)}$, $n = 1,2$ and C_{MX}^{ϕ}	
Electrolyte mixtures parameters superblock	c:
• Species triplets; ${}^{S}\theta_{MM'}$ and $\psi_{MM'X}$, or ${}^{S}\theta_{XX'}$ and $\psi_{MXX'}$	
Chemical elements block:	
• Elements, atomic weights	
Aqueous species superblock:	
Strict basis species	
• Auxiliary basis species, with log K	
on the standard grid	
• Non-basis species, with <i>log K</i>	
on the standard grid	
Pure minerals superblock:	
• Pure minerals, with <i>log K</i>	
on the standard grid	
Pure liquids superblock:	
• Pure liquids, with <i>log K</i>	
on the standard grid	
Gas species superblock:	
• Gas species, with log K on the standard	1
grid	
Solid solutions superblock:	
• Solid solutions	
References block:	
• References	

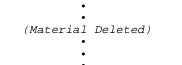
Figure 3. The basic structure of the data0 file for the hmw archetype.

The superblock for Pitzer parameters for pure aqueous neutral electrolytes is illustrated by the following, which is taken from **data0.hmw.R10**. This superblock begins with the header "sin-gle-salt parameters". The contents of a typical block, illustrated by that for the species pair "na+ cl-", is obvious. A block can also be entered for a species pair "so4-- co2(aq)". Note that here the "beta" parameters are really lambda parameters. EQPT maps the observable Pitzer parameters and their corresponding temperature derivatives in this superblock into corresponding conventional primitive equivalents. The conventional primitive equivalents are then written onto the **data1** file. The mapping relations are given in Chapter 3. Note that the standard temperature grid is not used to deal with the temperature dependence of any of the parameters of Pitzer's (1973, 1975, 1979, 1987) equations, with the exception of the A_{ϕ} Debye-Hückel parameter (which is not unique to these equations).

```
(a12)
single-salt parameters
             cl-
                                                                               (a12,2x,a12)
na+
               -1
                                                                               (f3.0,t15,f3.0)
  1
     beta0 = 0.07650 beta1 = 0.26440 beta2 = 0.00000 (2x,3(11x,f9.5)
alpha1 = 2.0 alpha2 = 12.0 (18x,2(16x,f5.1))
*
      cphi = 0.00127
                                                                            (13x, f9.5, 12x, f5.1)
    source = 84har/mol
                                                                               (13x,a18)
*
   db0/dt =0.000E+00d2b0/dt2 =0.000E+00db1/dt =0.000E+00d2b1/dt2 =0.000E+00db2/dt =0.000E+00d2b2/dt2 =0.000E+00dc/dt =0.000E+00d2c/dt2 =0.000E+00
                                                                          (13x,e10.3,13x,e10.3)
    source =
                                                                               (13x,a18)
                           _____
              (Material Deleted)
                                   _____
```

The superblock for Pitzer parameters for mixtures of two aqueous neutral electrolytes is illustrated by the following, which is taken from **data0.hmw.R10**. This superblock begins with the header "mixture term parameters". The contents of a typical block, illustrated by that for the species triplet "na+ k+ cl-", is obvious. EQPT maps the observable Pitzer parameters and their corresponding temperature derivatives in this superblock into corresponding conventional primitive equivalents. The conventional primitive equivalents are then written onto the **data1** file. The mapping relations are given in Chapter 3.

```
(a)
mixture term parameters
(a12,2(2x,a12))
na+
            k+
                 cl-
   theta = -0.01200 psi = -0.00180
source = 84har/mol
                                                            (13x, f8.5, 13x, f8.5)
                                                               (13x,a18)
*
*
  dth/dt = 0.0000 d2th/dt2 = 0.000E+00
dpsi/dt = 0.0000 d2ps/dt2 = 0.000E+00
                                                          (13x,e10.3,13x,e10.3)
   source =
                                                               (13x,a18)
```



+-----

4.3. The com Archetype for the data1/data1f Files

The **data1** file, being unformatted, can not be printed or displayed in meaningful fashion. The **data1f** file is a formatted equivalent. It can be useful in debugging if there are problems with a **data0** file or with EQPT. It also suffices to illustrate the structure of the **data1** file. In essence, each line of data in **data1f** represents one logical record of data in **data1**. Some block terminator lines appear in **data1f** to assist readability. The string "endit." is frequently used to mark the end of superblocks.

The **data1f** file corresponding to **data0.com.R10** is presented below. Material has been deleted where appropriate in order to present the essential facts, following the practice established earlier in presenting the major parts of the **data0** file structure. Some notes are superimposed on the material presented below. These are distinguished by the use of bold italic font.

Note that the structure of the **data1/data1f** files differs in some ways from that of the corresponding **data0** file. In particular, the superblocks containing activity coefficient data (here the "bdot" parameters) appear at the end of the file, instead of near the beginning. Also, the nominal temperature limits appear in a different location from that in which the data for the other "data0" parameters. The structure of the species blocks is also slightly different. EQPT computes the molecular weight of each species and puts this in the species block written on **data1/data1f**. Note that all data represented on the **data0** file on the standard temperature grid are replaced by the coefficients of interpolating polynomials. The first 5 coefficients represent a fit to the data in the range 0-100°C, the second five the data in the range 100-300°C. The first example of this in the material below occurs for the pressure "press".

A **data1/data1f** file begins with a record/line containing the string "data1". This is followed by one containing the key string "stfipc" (**com** archetype) or "stpitz" (**hmw** archetype). The third record/line contains the number of chemical elements and the number of basis species (strict plus auxiliary). The fourth record/line contains the name of the **data0** file used to generate the present **data1/data1f** file. This record/line is appended to the title as its first line.

dat THE	ipc 78 147 a0.com.R10 RMODYNAMIC DATABASE erated by gembochs/ING	(key string for a file of the c (number of chemical elements, n RES 15-apr-91	
+	0.0000 300.0000	(temperature limits)	
o ag al	15.99940 107.86820 26.98154	0.00000 0.00000 0.00000	(the zeros in the this block represent the oxide factor, which is no longer used)
	•		2

(Material Deleted) ٠ 91.22400 0.00000 zr press 1.013200000E+00 0.00000000E+00 0.00000000E+00 0.00000000E+00 0.00000000E+00 -4.345000000E-01 7.632333333E-03 5.514000000E-05-1.263733333E-06 1.396800000E-08 adh 4.938943704E-01 5.628991135E-04 5.680199734E-06-7.486279129E-09 0.00000000E+00 6.123000000E-01-2.559500000E-03 3.578166667E-05-1.378000000E-07 2.313333333E-10 bdh 3.252987270E-01 1.272991554E-04 5.443822909E-07-1.372465408E-09 0.00000000E+00 3.311000000E-01-2.133333333E-05 1.913333333E-06-7.0666666667E-09 1.0666666667E-11 bdot 1.738490207E-02 1.510054148E-03-2.607766840E-05 1.383863672E-07 0.00000000E+00 1.09000000E-01-1.483333333E-03 1.173333333E-05-3.4666666667E-08 2.666666667E-11 cco2 -1.031200000E+00 1.280600000E-03 2.559000000E+02 4.445000000E-01-1.606000000E-03 xlkeh -9.104826360E+01 3.463716921E-01-1.212214387E-03 2.669138257E-06 0.00000000E+00 -9.006110000E+01 3.154611667E-01-8.761516667E-04 1.533533333E-06-1.211333333E-09 aqueous 0 h2o 2 (name, # of elements, # of species in reaction) 18.015 Ο. (molecular weight, charge) 1.0000 o 2.0000 h (Material Deleted) 0 (name, # of elements, # of species in reaction) o2(g) 1 31.999 0. (molecular weight, charge) 2.0000 o (Material Deleted) (name, # of elements, # of species in reaction) acetic acid(aq) 3 4 60.053 0. (molecular weight, charge) 2.0000 2.0000 o С 4.0000 h -2.0000 o2(g) 2.0000 hco3--1.0000 acetic acid(aq) 2.0000 h+ 1.504679814E+02-6.181568055E-01 1.997116069E-03-4.266161503E-06 0.00000000E+00 1.486100000E+02-5.620795000E-01 1.405498333E-03-2.245400000E-06 1.4566666667E-09 (Material Deleted) (npo2)2(oh)2++ 4 (name, # of elements, # of species in reaction) 3 572.108 2. (molecular weight, charge) 2.0000 h 2.0000 np 6.0000 0 -1.0000 (npo2)2(oh)2++ -2.0000 h+ 2.0000 h2o 2.0000 npo2++ 7.128571429E+00-3.176190476E-02 1.047619048E-04 0.000000000E+00 0.00000000E+00 5.80000000E+00-8.00000000E-03 0.00000000E+00 0.00000000E+00 0.00000000E+00 (Material Deleted) ٠ •

endit. minerals (pb(oh)2)3.pbcl2(name, # of elements, # of species in reaction) 4 5 0. 0.000 1001.749 (molecular weight, charge, molar volume) 2.0000 cl 4.0000 pb 6.0000 o 6.0000 h -1.0000 (pb(oh)2)3.pbcl2 2.0000 cl--6.0000 h+ 4.0000 pb++ 6.0000 h2o 1.727930000E+01 0.00000000E+00 0.00000000E+00 0.00000000E+00 0.00000000E+00 5.00000000E+02 0.00000000E+00 0.0000000E+00 0.00000000E+00 0.0000000E+00 (Material Deleted) endit. gases 5 (name, # of elements, # of species in reaction) ag(g) 1 0.000 (molecular weight, charge, molar volume) 107.868 Ο. 1.0000 ag -1.0000 ag(g) -1.0000 h+ -0.2500 o2(g) 1.0000 ag+ 0.5000 h2o 5.554995563E+01-2.255708754E-01 7.834836108E-04-1.712704203E-06 0.00000000E+00 5.297980000E+01-1.615240000E-01 2.287600000E-04 0.00000000E+00 0.00000000E+00 (Material Deleted) endit. solid solutions (Material Deleted) (name, # of end members, # of site parameters) carbonate-calcite 1 6 1.000 calcite 1.000 magnesite 1.000 siderite 1.000 rhodochrosite 1.000 strontianite 1.000 smithsonite 0.000 0.000 0.000 0.000 0.000 0.000 1.000 0.000 0.000 0.000 0.000 0.000 (Material Deleted) endit. (uo2)3(co3)6(6-)4.0 0 (name, hard core diameter, insgfl flag) np(co3)5(6-) 4.0 0 u(co3)5(6-) 4.0 0 (Material Deleted) th6(oh)15(9+) 6.0 0 endit.

4.4. The hmw Archetype for the data1/data1f Files

The **data1f** file corresponding to **data0.hmw.R10** is presented below. Material has been deleted where appropriate in order to present the essential facts, following the practice established in the previous example. The superblocks containing the Pitzer coefficient parameters appear at the end of the file. Note that the observable parameters have been mapped into conventional primitive equivalents.

data1 stpitz (key string for a file of the com archetype) 9 13 (number of chemical elements, number of basis species) data0.hmw.R10 THERMODYNAMIC DATABASE generated by gembochs/INGRES 16-apr-91 _____ +---_ _ _ _ _ _ _ _ _ _ _ _ _ _____ 25.0000 25.0000 (temperature limits) +-----_____ (the zeros in the this block 0.00000 15.99940 0 0.00000 represent the oxide factor, which is no longer used) 40.07800 са 35.45270 0.00000 c] h 1.00794 0.00000 12.01100 0.00000 С k 39.09830 0.00000 mq 24.30500 0.00000 22.98977 0.00000 na s 32.06600 0.00000 press 1.013200000E+00 0.00000000E+00 0.00000000E+00 0.00000000E+00 0.00000000E+00 -4.345000000E-01 7.632333333E-03 5.514000000E-05-1.263733333E-06 1.396800000E-08 aphi 3.769945819E-01 5.417897788E-04 2.121889093E-06 8.607549305E-09 0.00000000E+00 4.540000000E-01-1.558333333E-03 2.361666667E-05-8.86666666667E-08 1.53333333E-10 xlkeh -9.104826360E+01 3.463716921E-01-1.212214387E-03 2.669138257E-06 0.00000000E+00 -9.006110000E+01 3.154611667E-01-8.761516667E-04 1.533533333E-06-1.211333333E-09 aqueous 2 0 (name, # of elements, # of species in reaction) h2o 18.015 0. (molecular weight, charge) 1.0000 o 2.0000 h (Material Deleted) o2(g) 1 0 (name, # of elements, # of species in reaction) 31.999 0. (molecular weight, charge) 2.0000 o (Material Deleted) caco3(aq) 3 4 (name, # of elements, # of species in reaction) 100.087 Ο. (molecular weight, charge) 1.0000 c 1.0000 ca 3.0000 0 -1.0000 caco3(aq) -1.0000 h+ 1.0000 ca++ 1.0000 hco3-7.188000000E+00 0.00000000E+00 0.00000000E+00 0.00000000E+00 0.00000000E+00 5.00000000E+02 0.00000000E+00 0.00000000E+00 0.00000000E+00 0.00000000E+00 (Material Deleted) .

endit. minerals (name, # of elements, # of species in reaction) anhydrite 3 3 0.000 136.142 0. (molecular weight, charge, molar volume) 1.0000 ca 1.0000 s 4.0000 o -1.0000 anhydrite 1.0000 so4--1.0000 ca++ -4.362100000E+00 0.000000000E+00 0.00000000E+00 0.00000000E+00 0.00000000E+00 5.00000000E+02 0.00000000E+00 0.0000000E+00 0.00000000E+00 0.0000000E+00 (Material Deleted) endit. gases 4 (name, # of elements, # of species in reaction) (molecular weight, charge molecular volume) co2(g) 2 44.010 0. 0.000 (molecular weight, charge, molar volume) 2.0000 o 1.0000 c -1.0000 co2(g) -1.0000 h2o 1.0000 h+ 1.0000 hco3--7.819200000E+00 0.00000000E+00 0.0000000E+00 0.00000000E+00 5.00000000E+02 0.00000000E+00 0.0000000E+00 0.0000000E+00 0.0000000E+00 (Material Deleted) endit. $(E_{\lambda} flag)$ E-lambda flag = on (off/on) Do not change this flag. na+ cl-+ cl-lambda0 = 0.07650 lambda1 = 0.26440 lambda2 = 0. alpha1 = 2.0 alpha2 = 12.0 dl0/dt = 0.000E+00 d210/dt2 = 0.000E+00 dl1/dt = 0.000E+00 d211/dt2 = 0.000E+00 dl2/dt = 0.000E+00 d212/dt2 = 0.000E+00 lambda2 = 0.00000 _____ +-----(Material Deleted) +-----_____ endit. cl-(species triplet for μ_{MMX} or μ_{MXX}) na+ na+ $mu = 0.00021 \ dmmx/dt = 0.000E+00 \ d2mmx/dt2 = 0.000E+00$ +-----(Material Deleted) _____ endit. na+ (species triplet for $\mu_{MM'X}$ or $\mu_{MXX'}$)
 na+
 cl (species triplet for

 mu = -0.00026
 dmu/dt = 0.000E+00
 d2mu/dt2 = 0.000E+00
 k+ _____ ٠ (Material Deleted)

endit. stop.

5. Code Architecture and Flow of Execution

The purpose of the present chapter is to provide a description of the structure of the software itself. This material is primarily included because it is required as part of the documentation to satisfy NUREG-0856 (Silling, 1983). It does not provide anything necessary for the typical code user. It may be helpful to those few users who desire to modify the code for whatever purpose.

In the present description, we will not make it a point to describe the role and function of every module in the source code. For such descriptions, the reader is referred to the relevant glossaries of modules. For EQPT modules, see Appendix B of the present report. For EQLIB modules, see Appendix A of the EQ3/6 Package Overview and Installation Guide (Wolery, 1992a). The purpose here is to describe the main features and essential aspects of the structure of the code. The purpose is not to provide detailed design documentation. Readers who want more detailed information are invited to examine the source code itself, which is reasonably well-documented internally.

The basic structure of EQPT is illustrated in the simplified flow diagram given in Figure 4. This diagram shows the flow of execution from the perspective of the main program, module **eqpt.f**. This is a fairly simple one-pass structure. EQPT does not process more than one **data0** file in a single run. There is branching in this flow only to deal with differences in the kinds of activity coefficient parameters and their processing between the **com** and **hmw** data file archetypes.

The simplified flow diagram omits references to writes to the **data1f**, **dpt1**, and **dpt2** files. The code writes to the **data1f** file whenever it writes to the **data1** file, as the former is a formatted version of the latter. The **dpt1** and **dpt2** files are only produced in the case of the **hmw** archetype. The **dpt1** file is written by modules **pdpz2.f** and **rdpz3.f**, and contains essentially just an echo of the observable Pitzer coefficient data read from the **data0** file. The **dpt2** file is written by module **wrpz3.f** and contains the corresponding conventional primitive Pitzer coefficient data. The same data are also written on the **data1f** file. The **dpt1** and **dpt2** files are vestigial in the present version of EQPT.

The simplified flow diagram also omits references to writes to the **slist** file and to some writes to the **output** and screen files which occur as the title and the superblocks for aqueous species, pure minerals, pure liquids, gas species, and solid solutions are processed by the relevant modules called by **eqpt.f**. These writes generate lists of the species on the data file. In the case of the screen file, only an abbreviated list is produced.

Module **eqpt.f** directs the overall process of code execution. Its first function is to get the time and date, which it does by calling the EQLIB module **timdat.f**. This module contains UNIX-dependent code. The main program then writes the code name and version identification, the copyright notice, and the time and date information to the screen and **output** files. Its next step is to initialize the dimensioning variables which correspond to the dimensioning parameters. Dimensioning variables are used to pass dimensioning data in the calling sequences of called modules; FORTRAN does not generally allow parameters (in the special FORTRAN sense) to be so passed.

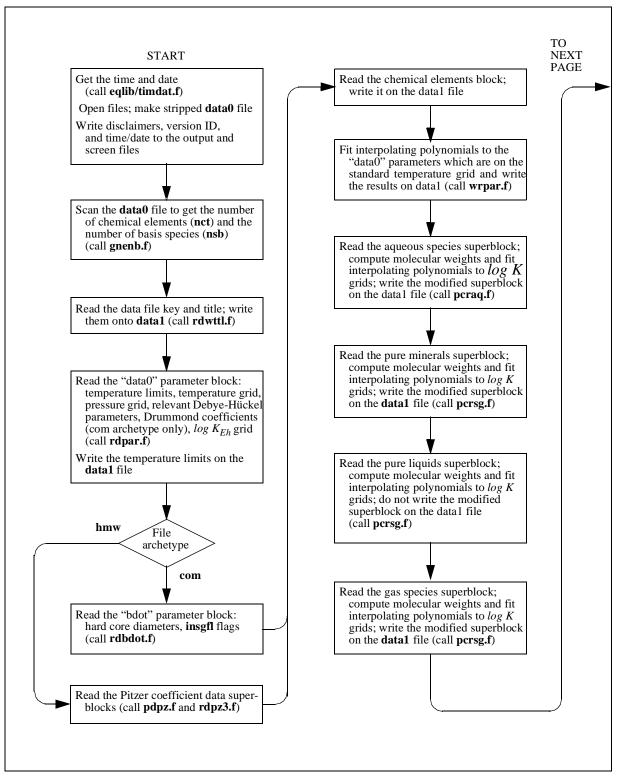


Figure 4. Simplified flow diagram of the EQPT code.

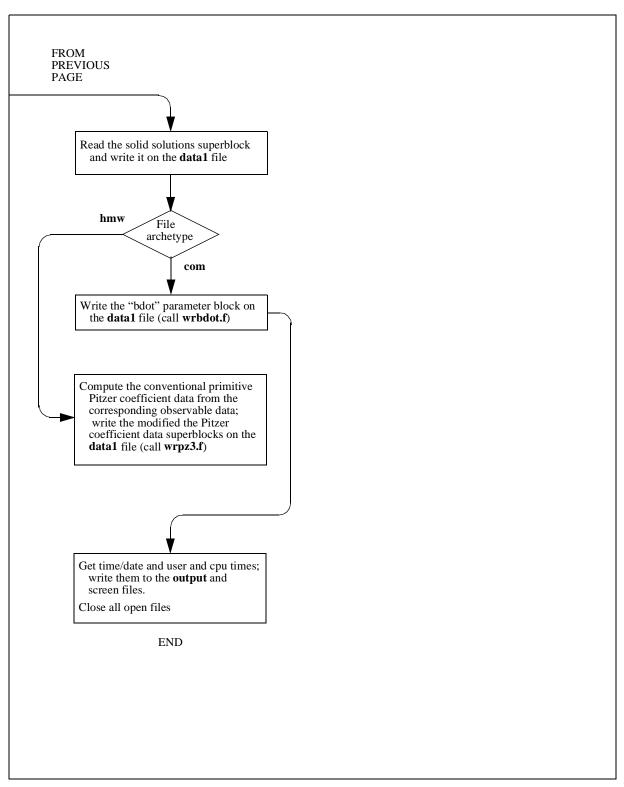


Figure 4 (continued). Simplified flow diagram of the EQPT code.

A seeming oddity in the flow of execution depicted in Figure 4 is that some block or superblocks are read from the **data0** file, processed, and written (sometimes in modified form) on the **data1** file in one continuous action, whereas in other cases some or all of the processing, and the writing of these blocks or superblocks on the **data1** file, does not occur until after succeeding blocks or superblocks have been read. This occurs because the succession of blocks and superblocks in the **data1** files differs somewhat from that of the corresponding **data0** files (see Chapter 4). There is no reason why this succession could not be the same for both kinds of files. However, if this were to be done, the activity coefficient superblocks should follow the species superblocks as in the **data1** files to facilitate processing of the data by EQ3NR and EQ6.

The main program module **eqpt.f** calls module **gnenb.f** to scan the **data0** file to determine the number of chemical elements (**nct**) and the number of basis species (**nsb**) on the data file. Module **eqpt.f** writes these data on the **data1** file, as well as the **output**, **slist**, and screen files.

Module **eqpt.f** calls module **rdwttl.f** to read the data file key (**com**, **nea**, **sup**, **hmw**, or **pit**) and the title. It validates the data file key against a set of known allowed possibilities and determines the code archetype. It then writes the data file key and the title on the **data1** file.

Module **eqpt.f** then calls module **rdpar.f** to read the "data0" parameter block. This includes the nominal temperature limits (°C) for the data file, the temperature values (°C) for the standard temperature grid, the pressure (bars) on the standard temperature grid, the relevant Debye-Hückel parameters on this grid ($A_{\gamma,10}$, B_{γ} and \dot{B} for the **com** archetype; A_{ϕ} for the hmw archetype), the coefficients for eq (12), which is used to compute the activity coefficient of aqueous CO_2 in *NaCl* solutions (**com** archetype only), and $log K_{Eh}$ on the standard temperature grid. The temperature limits are written on the **data1** file at this point. However, processing and writing of data for other parameters on this block is deferred (carried out later by module **wrpar.f**).

The activity coefficient data are then read. In the case of the **com** archetype, **eqpt.f** calls module **rdbdot.f** to read the block of "bdot" parameters (species names, hard core diameters, **insgfl** flags). In the case of the **hmw** archetype, the main program calls module **pdpz.f** to read the superblock of observable Pitzer parameters corresponding to solutions of pure aqueous neutral electrolytes (species pairs). It then calls module **rdpz.f** to read the superblock of observable Pitzer parameters of two aqueous neutral electrolytes.

Module **eqpt.f** then calls module **rdwele.f**. This reads the chemical elements block and writes it on the **data1** file. Following that, **eqpt.f** calls module **wrpar.f**. This module processes the remaining "data0" parameters and writes the corresponding data on the **data1** file. The parameters which were read from the **data0** file in the form of values on the standard temperature grid are processed as follows Module **wrpar.f** calls module **intrp.f** in each case to fit one interpolating polynomial to the part of the grid for the temperature range 0-100°C, and a second such polynomial to the part of the grid for the range 100-300°C. The interpolating polynomial coefficients replace the original grid values in the data written onto the **data1** file.

Module **eqpt.f** then processes four species superblocks. The aqueous species superblock is processed by a call to module **pcraq.f**. This module reads a data block for each species in the superblock, processes it, and writes a corresponding block onto the **data1** file. The *log K* data for the associated reaction (all aqueous species but strict basis species have such a reaction) are read

from the **data0** file as values on the standard temperature grid. Module **pcraq.f** calls module **rxnchk.f** to check each associated reaction for mass and charge balance. Module **pcraq.f** then calls module **intrp.f** to fit interpolating polynomials to these data in the same manner as it does for the "data0" parameters which are read as values on this grid. When the data block for the species is written onto the **data1** file, the interpolating polynomial coefficients replace the values on the grid. Module **pcraq.f** also computes the molecular weight of each species and includes this in the data block written on the **data1** file.

The superblocks for pure minerals, pure liquids, and gas species are handled similarly by a call in each case to module **pcrsg.f**. The pure liquids superblock is read, but this superblock is not written onto the **data1** file, as EQ3NR and EQ6 have not been developed to deal with pure non-aqueous liquids. Module **pcrsg.f** is closely analogous to module **pcraq.f**.

Module **eqpt.f** then calls module **pcrss.f**. This reads the solid solutions superblock and writes it onto the **data1** file. This superblock is not analogous to those for aqueous species, pure minerals, pure liquids, or gas species, as solid solutions are *phases*, not species. The data block for each solid solution includes a list of end-member components, which must correspond to pure minerals appearing in the pure minerals superblock.

The activity coefficient data are then processed and written to the **data1** file. In the case of the **com** archetype, the data require no processing. Module **eqpt.f** calls module **wrbdot.f** to write the "bdot" block onto the **data1** file. In the case of the **hmw** archetype, **eqpt.f** calls module **wrpz3.f**. This module processes the data, calculating conventional primitive Pitzer coefficient data from the set of corresponding observable coefficient data read from the **data0** file. Three superblocks of conventional primitive coefficient data are then written on the **data1** file: one for the λ coefficients corresponding to all types of species pairs, one for μ coefficients for species triplets in which no species appears twice, and one for μ coefficients for species triplets in which no species appears more than once. This contrasts with the two superblocks of observable coefficient data that are read from the **data0** file.

All data file processing is now complete. Module **eqpt.f** gets the time and date and user and cpu times and writes them to the **output** and screen files. It then closes all open files and terminates execution.

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Appendix A: Glossary of Major Variables in EQPT

This glossary covers the major variables in EQPT. Most all of these are shared with EQ3NR (Wolery, 1992b) and EQ6 (Wolery and Daveler, 1992). Those which are unique to EQPT are so marked. This glossary does not include variables which are of a purely local nature, such as a floating point variable used to accumulate sums in a do loop. Logarithmic quantities are frequently used. All refer to base ten, unless otherwise specified.

The variable names may be preceded by the corresponding algebraic symbols used in this report, if any. Following the conventions used universally in the EQ3/6 package, variables beginning with **i**, **j**, **k** or **n** are integer, those beginning with **q** are logical, and those beginning with **u** are character variables. Variables beginning with any other letters, including **l** and **m**, should be **re-al*8**. On 32-bit machines, this corresponds to **double precision**; on 64-bit machines, this corresponds to **single precision**.

Most arrays in EQ3/6 are dimensioned using FORTRAN parameters. The value of a parameter is assigned once in a code (in a PARAMETER statement) and can not be changed elsewhere. Some FORTRAN compilers permit parameters to be passed through calling sequences, but others do not allow this, as a means of protecting the parameter's assigned value. A called module often requires the dimensions of passed arrays. This is handled in EQ3/6 by the use of "dimensioning variables," which are ordinary integer variables whose values have been set equal to the corresponding parameters. The names of the dimensioning variables end, respectively, in **par**, **pa**, or **pa1**. The names of the corresponding dimensioning variables end, respectively, in **max**, **mx**, and **mx1**. Thus, the dimension of the **uelem** matrix is given by the pair **nctpar/nctmax** (the maximum number of chemical elements. The parameter suffixes are used exclusively by dimensioning variables. For example, **itermx** is the number of Newton-Raphson iterations in EQ3/R and EQ6.

Α _{γ,10}	adh	Debye-Hückel A_{γ} parameter for computing the base ten logarithm of the molal ac- tivity coefficient of an aqueous species. Dimensioning: adh(8) . Usage: adh(n) is the Debye-Hückel A_{γ} parameter for the n -th point on the standard temperature grid. Note: dimensioning and usage differ from that in EQ3NR and EQ6.
α ₁	alph1	The Pitzer α_I parameter. Dimensioning: alph1(npxpar) . Usage: alph1(npxp) refers to the α_I parameter.for the npxp -th neutral electrolyte (cation-anion pair). The arrays alph1 and alph2 are equivalent to the palpha array in EQ3NR and EQ6.
α ₂	alph2	The Pitzer α_2 parameter. Dimensioning: alph2(npxpar) . Usage: alph2(npxp) refers to the α_2 parameter.for the npxp -th neutral electrolyte (cation-anion pair). The arrays alph1 and alph2 are equivalent to the palpha array in EQ3NR and EQ6.
$A_{igoplus}$	aphi	Debye-Hückel A_{ϕ} parameter. Dimensioning: aphi(8) . Usage: aphi(n) is the Debye-Hückel A_{ϕ} parameter for the n -th point on the standard temperature grid. Note: dimensioning and usage differ from that in EQ3NR and EQ6.
	apr	Work array for computing interpolating polynomial coefficients to fit the xlks array. Dimensioning: apr(10). Usage: apr(i) refers to the i -th coefficient for the current

		species. The apr array is copied into the part of the ars array corresponding to the current species. This array is unique to EQPT.
	арх	Solid solution activity coefficient model parameters. Dimensioning: apx(12). Usage: apx(i) refers to the i -th parameter for the current solid solution. Note: dimensioning and usage differ from that in EQ3NR and EQ6.
	ars	Interpolating polynomial coefficients for computing the array of equilibrium con- stants xlks . Dimensioning: ars(10,nsqpa1) . Usage: ars(i,ns) refers to the i -th coef- ficient for the ns -th species. Note: dimensioning and usage differ from that in EQ3NR and EQ6.
${}^{S}\theta_{MN}, {}^{S}\lambda_{MN}^{(0)}$	atheta	The Pitzer short range theta coefficient (25°C). Dimensioning: atheta(npxpar). Us-
		age: atheta (k) refers to the ${}^{S}\theta$ coefficient k -th cation-cation or anion-anion pair.
		Technically, this is the average ${}^{S}\theta$ value, as a different value may be fit to more than one mixture of two aqueous neutral electrolytes. Individual values for such mixtures are read from the data0 file into the theta array, and atheta is computed from them. However, a common value should be specified on the data file for all such mixtures.
		This array is unique to EQPT. The coefficient ${}^{S}\theta_{MN}$ is mapped to the primitive Pitzer
		coefficient ${}^{S}\lambda_{MN}$ (the two are equal by the convention used in EQ3/6). There is no
		ionic strength dependence for this particular λ_{ij} coefficient, so ${}^{S}\lambda_{MN} = {}^{S}\lambda_{MN}^{(0)}$. In
		EQ3NR and EQ6, these and other $\lambda_{ij}^{(n)}$ coefficients are read into the bslm array.
	atwt	Atomic weight of a chemical element. Dimensioning: atwt(nctpar) . Usage: atwt(nc) refers to the nc -th chemical element.
B_{γ}	bdh	Debye-Hückel B_{γ} parameter. Dimensioning: bdh(8) . Usage: bdh(n) is the Debye-Hückel B_{γ} parameter for the n -th point on the standard temperature grid. Note: dimensioning and usage differ from that in EQ3NR and EQ6.
<i>B</i>	bdot	Extended Debye-Hückel \dot{B} parameter. Dimensioning: bdot(8) . Usage: bdot(n) is
		the extended Debye-Hückel \dot{B} parameter for the n -th point on the standard temper- ature grid. Note: dimensioning and usage differ from that in EQ3NR and EQ6.
	cco2	Parameters of the Drummond (1981) equation for computing $\log \gamma_{CO_{2(aq)}}$ as a func-
		tion of temperature and ionic strength. Dimensioning: $cco2(5)$. Usage: $cco2(i)$ refers to the i -th coefficient.
b _{sr}	cdrs	Reaction coefficient array. Dimensioning: cdrs(nsqpa2,nsqpa1). Usage:
		cdrs (nse , ns) is the coefficient of the nse -th aqueous basis species appearing in the reaction for the ns -th species; cdrg (nsq1 , ns) is the coefficient of the ns -th species itself. If the ns -th species is an auxiliary basis species then cdrs (nsq1 , ns) = 0 and cdrs (nsq1 , nrs) is the corresponding coefficient. Note: dimensioning and usage differ from that in EQ3NR and EQ6. The cdrs array of EQPT encompasses the cdrs , cdrm , and cdrg arrays of EQ3NR and EQ6.
	cdum1	Holding array used to read in the elemental composition coefficients of species list- ed on the data file. Dimensioning: cdum1(nsqpa1) . Usage: cdum1(n) is the n -th co- efficient for the current species as it is written on the data file; it corresponds to the

element whose name is **unam8(n)**. This array is known as **cessd** in EQ3NR and EQ6.

 $c_{\varepsilon s}$ **cess** Array containing the elemental composition coefficients of aqueous species. Dimensioning: **cess(nctpar,nsqpa1)**. Usage: **cess(nc,ns)** is the coefficient of the **nc**-th chemical element for the **ns**-th aqueous species. Note: dimensioning and usage differ from that in EQ3NR and EQ6. The **cess** array of EQPT encompasses the **cess, cemn**, and **cegs** arrays of EQ3NR and EQ6.

cph The Pitzer coefficient C_{MX}^{ϕ} (25°C). Dimensioning: **cph**(**npxpar**). Usage:

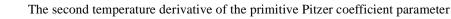
cph(npxp) is the C^{ϕ} coefficient for the **npxp**-th pure aqueous neutral electrolyte. This array is unique to EQPT. This coefficient is used by EQPT to compute the primitive Pitzer coefficients μ_{MMX} and μ_{MXX} , which are respectively stored in the **mum-mx** and **mumxx** arrays. In EQ3NR and EQ6, these and other μ_{ijk} coefficients are read into the **bmu** array.

The second temperature derivative of the Pitzer coefficient C_{MX}^{ϕ} (25°C). Dimen-

sioning: **d2c(npxpar)**. Usage: **d2c(npxp)** is the second temperature derivative of the C^{ϕ} coefficient for the **npxp**-th pure aqueous neutral electrolyte. This array is unique to EQPT, which uses. $d^2 C^{\phi}_{MX}/dT^2$ to compute $d^2 \mu_{MMX}/dT^2$ and $d^2 \mu_{MXX}/dT^2$, which are respectively stored in the **d2mmx** and **d2mxx** arrays In EQ3NR and EQ6, these and corresponding derivatives of other μ_{ijk} coefficients are read into the **dmu2** array.

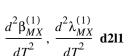
The second temperature derivative of the primitive Pitzer coefficient parameter

 $\lambda_{MX}^{(0)}$ (25°C values). By convention, $\lambda_{MX}^{(0)}$ equals the Pitzer coefficient parameter $\beta_{MX}^{(0)}$, so the derivatives are also equal to one another. Dimensioning: **d2l0(npxpar)**. Usage: **d2l0(npxp)** is the second derivative for the **npxp**-th aqueous neutral electrolyte. This array is unique to EQPT. In EQ3NR and EQ6, these and the corresponding derivatives of other $\lambda_{ij}^{(n)}$ parameters are read into the **dslm2** array.



 $\lambda_{MX}^{(1)}$ (25°C values). By convention, $\lambda_{MX}^{(1)}$ equals the Pitzer coefficient parameter

 $\beta_{MX}^{(1)}$, so the derivatives are also equal to one another. Dimensioning: **d2l1(npxpar)**. Usage: **d2l1(npxp)** is the second derivative for the **npxp**-th aqueous neutral electrolyte. This array is unique to EQPT. In EQ3NR and EQ6, these and the corresponding derivatives of other $\lambda_{ii}^{(n)}$ parameters are read into the **dslm2** array.



 $\frac{d^2 \beta_{MX}^{(0)}}{dT^2}, \frac{d^2 \lambda_{MX}^{(0)}}{dT^2}$ d2l0

d2c

 C_{MX}^{ϕ}



$\frac{d^2 \beta_{MX}^{(2)}}{dT^2}, \frac{d^2 \lambda_{MX}^{(2)}}{dT^2}$	d2l2	The second temperature derivative of the primitive Pitzer coefficient parameter
al al		$\lambda_{MX}^{(2)}$ (25°C values). By convention, $\lambda_{MX}^{(2)}$ equals the Pitzer coefficient parameter
		$\beta_{MX}^{(2)}$, so the derivatives are also equal to one another. Dimensioning: d2l2(npxpar) .
		Usage: d2l2(npxp) is the second derivative for the npxp -th aqueous neutral electro- lyte. This array is unique to EQPT. In EQ3NR and EQ6, these and the corresponding
		derivatives of other $\lambda_{ii}^{(n)}$ parameters are read into the dslm2 array.
2		
$\frac{d^2\mu_{MMX}}{dT^2}$	d2mmx	The second temperature derivative of the primitive Pitzer coefficient μ_{MMX} (25°C).
		Dimensioning: d2mmx(npxpar). Usage: d2mmx(npxp) is the second temperature derivative of the μ_{MMX} coefficient for the npxp -th pure aqueous neutral electrolyte. It is computed from the d2c array. This array is unique to EQPT. In EQ3NR and EQ6, these and corresponding derivatives of other μ_{ijk} coefficients are read into the dmu2 array.
$\frac{d^2\mu_{MNX}}{dT^2}$	d2mu	The second temperature derivative of the primitive Pitzer coefficient μ_{MXX} (25°C).
<i>u</i> 1		This is a local variable in module rdpz3.f computed from the d2psi , d2mmx , and d2mxx arrays. In EQ3NR and EQ6, these and corresponding derivatives of other μ_{ijk} coefficients are read into the dmu2 array.
$\frac{d^2\mu_{MXX}}{dT^2}$	d2mxx	The second temperature derivative of the primitive Pitzer coefficient μ_{MXX} (25°C).
u I		Dimensioning: d2mxx(npxpar) . Usage: d2mxx(npxp) is the second temperature derivative of the μ_{MXX} coefficient for the npxp -th pure aqueous neutral electrolyte. It is computed from the d2c array. This array is unique to EQPT. In EQ3NR and EQ6, these and corresponding derivatives of other μ_{ijk} coefficients are read into the dmu2 array.
$\frac{d^2 \Psi_{MNX}}{dT^2}$	d2psi	The second temperature derivative of the Pitzer coefficient ψ_{MNXZ} (25°C). Dimen-
<i>a</i> 1		sioning: d2psi(npxpar). Usage: d2psi(nmpp) is the second temperature derivative of the ψ coefficient for the nmpp -th mixture of two aqueous neutral electrolytes containing a common ion <i>X</i> . This array is unique to EQPT, which uses. $d^2 \psi_{MNX}/dT^2$ to compute $d^2 \mu_{MNX}/dT^2$ and $d^2 \mu_{MNX}/dT^2$, which are respectively stored in the d2mu variable. In EQ3NR and EQ6, these and corresponding derivatives of other μ_{ijk} coefficients are read into the dmu1 array.
$\frac{d^{2S}\theta_{MN}}{dT^2}, \frac{d^2\lambda_M^{(0)}}{dT^2}$)) <u>N</u>	



The second temperature derivative of the Pitzer short range theta coefficient (25° C). Dimensioning: **d2th**(**npxpar**). Usage: **d2th**(**k**) refers to the second temperature de-

rivative of the ^S θ coefficient for the **k**-th cation-cation or anion-anion pair. Technically, this is an average value, as a different value may be fit to more than one mixture of two aqueous neutral electrolytes. Individual values for such mixtures are read from the **data0** file into the **d2thd** variable for each mixture, and **d2th** is computed as the average value. However, a common value should be specified on the data file for all such mixtures. This array is unique to EQPT. The coefficient $d^{2S}\theta_{MN}/dT^2$ is mapped to the primitive Pitzer coefficient $d^{2S}\lambda_{MN}/T^2$ (the two are equal by the convention used in EQ3/6). There is no ionic strength dependence for this particular coefficient, so $d^{2S}\lambda_{MN}/dT^2 = d^2\lambda_{MN}^{(0)}/dT^2$. In EQ3NR and EQ6, these and corresponding derivatives of the $\lambda_{ij}^{(n)}$ parameters are read into the **dsIm2** array.

$\frac{d^{2S}\theta_{MN}}{dT^2}$	d2thd	The second temperature derivative of the Pitzer short range theta coefficient (25°C). This is a local variable in module rdpz3.f used to read in the value of this derivative for the current mixture of two aqueous neutral electrolytes containing a common ion. See d2thd .
$\frac{dC^{\phi}_{MX}}{dT}$	dc	The first temperature derivative of the Pitzer coefficient C_{MX}^{ϕ} (25°C). Dimension- ing: dc(npxpar) . Usage: dc(npxp) is the first temperature derivative of the C^{ϕ} coef- ficient for the npxp -th neutral aqueous neutral electrolyte. This array is unique to
		EQPT, which uses. dC_{MX}^{ϕ}/dT to compute $d\mu_{MMX}/dT$ and $d\mu_{MXX}/dT$, which are respectively stored in the dmmx and dmxx arrays. In EQ3NR and EQ6, these and corresponding derivatives of other μ_{ijk} coefficients are read into the dmu1 array.
$\frac{d\beta_{MX}^{(0)}}{dT}, \frac{d\lambda_{MX}^{(0)}}{dT}$	d10	The first temperature derivative of the primitive Pitzer coefficient parameter $\lambda_{MX}^{(0)}$, (25°C values). By convention, $\lambda_{MX}^{(0)}$ equals the Pitzer coefficient parameter $\beta_{MX}^{(0)}$, so the derivatives are also equal to one another. Dimensioning: dl0(npxpar) . Usage: dl0(npxp) is the first derivative for the npxp -th aqueous neutral electrolyte. This array is unique to EQPT. In EQ3NR and EQ6, these and the corresponding derivatives of other $\lambda_{ii}^{(n)}$ parameters are read into the dslm1 array.
$\frac{d\beta_{MX}^{(1)}}{dT}, \frac{d\lambda_{MX}^{(1)}}{dT}$	d11	The first temperature derivative of the primitive Pitzer coefficient parameter $\lambda_{MX}^{(1)}$ (25°C values). By convention, $\lambda_{MX}^{(1)}$ equals the Pitzer coefficient parameter $\beta_{MX}^{(1)}$, so the derivatives are also equal to one another. Dimensioning: dl1(npxpar) . Usage: dl1(npxp) is the first derivative for the npxp -th aqueous neutral electrolyte. This array is unique to EQPT. In EQ3NR and EQ6, these and the corresponding derivatives of other $\lambda_{ij}^{(n)}$ parameters are read into the dslm1 array.

$\frac{d\beta_{MX}^{(2)}}{dT}, \frac{d\lambda_{MX}^{(2)}}{dT}$	d12	The first temperature derivative of the primitive Pitzer coefficient parameter $\lambda_{MX}^{(2)}$
		(25°C values). By convention, $\lambda_{MX}^{(2)}$ equals the Pitzer coefficient parameter $\beta_{MX}^{(2)}$, so
		the derivatives are also equal to one another. Dimensioning: dl2(npxpar) . Usage: dl2(npxp) is the first derivative for the npxp -th aqueous neutral electrolyte. This array is unique to EQPT. In EQ3NR and EQ6, these and the corresponding derivatives
		of other $\lambda_{ij}^{(n)}$ parameters are read into the dslm1 array.
$\frac{d\mu_{MMX}}{dT}$	dmmx	The first temperature derivative of the primitive Pitzer coefficient μ_{MMX} (25°C). Di-
		mensioning: dmmx(npxpar) . Usage: dmmx(npxp) is the first temperature deriva- tive of the μ_{MMX} coefficient for the npxp -th pure aqueous neutral electrolyte. It is computed from the dc array. This array is unique to EQPT. In EQ3NR and EQ6, these and corresponding derivatives of other μ_{ijk} coefficients are read into the dmu1 array.
$\frac{d\mu_{MNX}}{dT}$	dmu	The first temperature derivative of the primitive Pitzer coefficient μ_{MXX} (25°C).
dT		This is a local variable in module rdpz3.f computed from the dpsi , dmmx , and dmxx arrays. In EQ3NR and EQ6, these and corresponding derivatives of other μ_{ijk} coefficients are read into the dmu1 array.
$\frac{d\mu_{MXX}}{dT}$	dmxx	The first temperature derivative of the primitive Pitzer coefficient μ_{MXX} (25°C). Di-
		mensioning: dmxx (npxpar). Usage: dmxx (npxp) is the first temperature derivative of the μ_{MXX} coefficient for the npxp -th pure aqueous neutral electrolyte. It is computed from the dc array. This array is unique to EQPT. In EQ3NR and EQ6, these and corresponding derivatives of other μ_{ijk} coefficients are read into the dmu1 array.
$rac{d\psi_{MNX}}{dT}$	dpsi	The first temperature derivative of the Pitzer coefficient ψ_{MNXZ} (25°C). Dimen-
		sioning: dpsi(npxpar) . Usage: dpsi(nmpp) is the first temperature derivative of the ψ coefficient for the nmpp -th mixture of two aqueous neutral electrolytes containing a common ion <i>X</i> . This array is unique to EQPT, which uses. $d\psi_{MNX}/dT$ to com-
		pute $d\mu_{MNX}/dT$ and $d\mu_{MNX}/dT$, which are respectively stored in the dmu variable. In EQ3NR and EQ6, these and corresponding derivatives of other μ_{ijk} coefficients are read into the dmu1 array.
$\frac{d^{S}\theta_{MN}}{dT}, \frac{d\lambda_{MN}^{(0)}}{dT}$		

dth

The first temperature derivative of the Pitzer short range theta coefficient (25°C). Dimensioning: **dth(npxpar)**. Usage: **dth(k)** is the first temperature derivative of the

 ${}^{S}\theta$ coefficient for the **k**-th cation-cation or anion-anion pair. Technically, this is an average value, as a different value may be fit to more than one mixture of two neutral aqueous neutral electrolytes. Individual values for such mixtures are read from the

data0 file into the **dthd** variable for each mixture, and **dth** is computed as the average value. However, a common value should be specified on the data file for all such mixtures. This array is unique to EQPT. The coefficient $d^S \theta_{MN}/dT$ is mapped to the primitive Pitzer coefficient $d^S \lambda_{MN}/T$ (the two are equal by the convention used in EQ3/6). There is no ionic strength dependence for this particular coefficient, so $d^S \lambda_{MN}/dT = d \lambda_{MN}^{(0)}/dT$. In EQ3NR and EQ6, these and corresponding derivatives of the $\lambda_{ii}^{(n)}$ parameters are read into the **dslm1** array.

dthd The first temperature derivative of the Pitzer short range theta coefficient (25° C). This is a local variable in module **rdpz3.f** used to read in the value of this derivative for the current mixture of two aqueous neutral electrolytes containing a common ion. See dth. eps100 One hundred times the real*8 machine epsilon. gdum A work array into which is copied the portion of the **xlks** array for a given species. Dimensioning: gdum(8). Usage: gdum(n) = xlks(n,ns) for the species currently indicated by ns. This array is local to modules pcraq.f and pcrsg.f. ier An error flag parameter commonly found in subroutine calling sequences. Values greater than zero mark error conditions. Values less than zero mark warning conditions. The significance of this flag is only that the subroutine did or may have failed to carry out its primary function. For example, a matrix solver may have found a matrix to be computationally singular. This situation may or may not equate to an error in the context of the larger code. For example, the larger code may not have a problem with the fact that a matrix is computationally singular. iktmax The maximum number of end members in a solid solution. This is the variable which corresponds to the dimensioning parameter iktpar. iktpar Dimensioning parameter: the maximum number of end members in a solid solution. See iktmax. iline The maximum number of characters per line in the data file. This is currently set to 80. ineu Flag variable for determining how to compute the activity coefficient of a neutral solute species in conjunction with the B-dot equation. This is a local variable in module wrbdot.f. Usage: ineu is the flag for the current aqueous species: = 0 Set $log \gamma_i = 0$ (polar species) = 1 Use the Drummond (1981) polynomial (non-polar species). This flag variable is stored in the insgfl array in EQ3NR and EQ6. insgfl See ineu. ixs The number of distinct Pitzer theta values read from the data file for a given pair of cations or anions. Dimension: ixs(npxpar). Usage: ixs(nmx) is the number of such values for the nmx-th such pair.

 $\frac{d^3 \theta_{MN}}{dT}$

$eta_{MX}^{(0)},\lambda_{MX}^{(0)}$	10	The primitive Pitzer coefficient parameter $\lambda_{MX}^{(0)}$, taken by convention to be equal to
		the Pitzer coefficient parameter $\beta_{MX}^{(0)}$ (25°C values). Dimensioning: 10(npxpar) . Usage: 10(npxp) is the parameter for the npxp -th aqueous neutral electrolyte. This
		array is unique to EQPT. In EQ3NR and EQ6, these and other $\lambda_{ij}^{(n)}$ parameters are read into the bslm array.
$eta_{MX}^{(1)},\lambda_{MX}^{(1)}$	11	The primitive Pitzer coefficient parameter $\lambda_{MX}^{(1)}$, taken by convention to be equal to
		the Pitzer coefficient parameter $\beta_{MX}^{(1)}$ (25°C values). Dimensioning: l1(npxpar) . Usage: l1(npxp) is the parameter for the npxp -th aqueous neutral electrolyte. This
		array is unique to EQPT. In EQ3NR and EQ6, these and other $\lambda_{ij}^{(n)}$ parameters are read into the bslm array.
$eta_{MX}^{(2)},\lambda_{MX}^{(2)}$	12	The primitive Pitzer coefficient parameter $\lambda_{MX}^{(2)}$, taken by convention to be equal to
		the Pitzer coefficient parameter $\beta_{MX}^{(2)}$ (25°C values). Dimensioning: 12(npxpar) .
		Usage: $l2(npxp)$ is the parameter for the npxp -th aqueous neutral electrolyte. This
		array is unique to EQPT. In EQ3NR and EQ6, these and other $\lambda_{ij}^{(n)}$ parameters are
		read into the bslm array.
μ_{MMX}	mummx	The primitive Pitzer coefficient μ_{MMX} (25°C). Dimensioning: mummx (npxpar).
		Usage: mummx(npxp) is the μ_{MMX} coefficient for the npxp -th pure aqueous neutral
		electrolyte. It is computed from the C^{ϕ} coefficient. See cph . This array is unique to EQPT. In EQ3NR and EQ6, these and other μ_{ijk} coefficients are read into the bmu array.
μ_{MNX}	mumnx	The primitive Pitzer coefficient μ_{MNX} (25°C). This is a local variable in module
• 10/10/A		rdpz3.f computed from the ψ coefficient. See psi . In EQ3NR and EQ6, these and other μ_{ijk} coefficients are read into the bmu array.
μ_{MXX}	mumxx	The primitive Pitzer coefficient μ_{MXX} (25°C). Dimensioning: mumxx(npxpar) . Us-
		age: mumxx(npxp) is the μ_{MXX} coefficient for the npxp-th pure aqueous neutral
		electrolyte. It is computed from the C^{ϕ} coefficient. See cph . This array is unique to EQPT. In EQ3NR and EQ6, these and other μ_{ijk} coefficients are read into the bmu array.
M _i	mwtss	Molecular weight of a chemical species. This is a local variable in modules pcraq.f and pcrsg.f . The dimensioning and usage differ in EQ3NR and EQ6.
ϵ_T	nct	Total number of chemical elements.
	nctmax	The maximum number of chemical elements. This is a variable which corresponds to the parameter nctpar .
	nctpa1	Dimensioning parameter: the maximum number of chemical elements + 1. See nctpar. This is presently unused.

nctpar	Dimensioning parameter: the maximum number of chemical elements. See nctmax .
ncts	The number of chemical elements comprising a chemical species. This is a local variable in modules pcraq.f and pcrsg.f . It is unique to EQPT.
ndata0	The unit number of the stripped data0 file. This is a scratch file that has been stripped of any comment lines. This variable is unique to EQPT.
ndata1	The unit number of the data1 file. This variable is unique to EQPT. In EQ3NR and EQ6, the unit number of this file is nad1 .
ndat1f	The unit number of the data1f file. This variable is unique to EQPT.
ndpt1	The unit number of the dpt1 file. This variable is unique to EQPT.
ndpt2	The unit number of the dpt2 file. This variable is unique to EQPT.
ndrs	The number of species in a chemical reaction. This is a local variable in modules pcraq.f and pcrsg.f . It is unique to EQPT.
nmpp	The number of mixtures of two aqueous neutral electrolytes for which Pitzer coef- ficients have been read from the data file. This variable is unique to EQPT.
nmtp	The number of distinct cation-cation or anion-anion pairs corresponding to Pitzer coefficients for mixtures of aqueous neutral electrolytes containing a common (third) ion. This is used in analyzing data preparatory to computing the average Pitzer theta coefficient for such pairs of ions. See atheta . This variable is unique to EQPT.
nout	The unit number of the out file. This variable and the corresponding file are no long- er used.
noutpt	The unit number of the output file.
прхр	Counter for the number of blocks of Pitzer coefficient data read from either the su- perblock for pure aqueous neutral electrolytes or the superblock for mixtures of two such electrolytes.
npxpar	Dimensioning parameter: the maximum number entries in the various Pitzer coefficient data arrays.
nsb	The number of strict basis species. Also the index denoting the aqueous redox species (currently O_2); nsb = nct + 1.
nslist	The unit number of the slist file. This variable is unique to EQPT.
nslt	The number of pure aqueous neutral electrolytes for which Pitzer coefficients have been read from the data file. This variable is unique to EQPT.
nsq	Number of aqueous basis species.
nsq1	The variable equivalent to $nsq + 1$. This is used in reaction coefficient arrays to mark the coefficient of the species formally associated with a given reaction. For example, cdrs(nsq1,ns) refers to the coefficient of the associated aqueous species destroyed

 s_B

 s_Q

in the reaction for the **ns**-th species. The first **nsq** species are the aqueous basis species. The **nsq1**-th species is the current non-basis species, which may be a species of any type of phase.

- nsq2 The variable equivalent to nsq + 2. This is unique to EQPT and is presently not used.
- **nsqmax** The maximum number of aqueous basis species. This is the variable corresponding to the parameter **nsqpar**.
- **nsqpa1** Dimensioning parameter: the equivalent of **nsqpar** + 1.
- **nsqpa2** Dimensioning parameter: the equivalent of **nsqpar** + 2.
- **nsqpar** Dimensioning parameter: the maximum number of aqueous basis species. See **nsqmax**.
- **nssp** The number of aqueous species appearing in the set of pure aqueous neutral electrolytes for which Pitzer coefficients have been read from the data file. This variable is unique to EQPT. See **ussp**.
- **nstpar** Dimensioning parameter: the maximum number of aqueous species.
- **ntitld** The number of lines of the title on the **data1** file.
- **ntitpa** Dimensioning parameter: the maximum number of lines in a data file title.
- ntmps The unit number of the data0 file. This variable is unique to EQPT.
- **nttyi** The unit number of the keyboard file. This variable is unique to EQPT and is currently not used.
- **nttyo** The unit number of the screen file.

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- **nubdot** The number of aqueous species for which hard core diameters and **insgfl** (**ineu**) flags are specified on the data file. See **ubdotp**.
- oxfacStoichiometric oxide factor. Dimensioning: oxfac(nctpar). Usage: oxfac(nc) is the
oxide factor for the nc-th chemical element. See uoxide. Both arrays are currently
vestigial, though they are processed by EQPT.
- pressPressure, bars, on the standard temperature grid. Dimensioning: press(8). Usage:
press(n) is the pressure for the n-th point on the grid. Note: dimensioning and usage
differ from that in EQ3NR and EQ6.
- Ψ_{MNX} **psi** The Pitzer coefficient Ψ_{MNX} (25°C). Dimensioning: **psi(npxpar**). Usage: **psi(nmpp)** is the Ψ coefficient for the **nmpp**-th mixture of two aqueous neutral electrolytes containing a common ion X. This array is unique to EQPT. This coefficient is used by EQPT to compute the primitive Pitzer coefficient μ_{MNX} , which is stored in the **mumnx** variable. In EQ3NR and EQ6, these and other μ_{ijk} coefficients are read into the **bmu** array.
 - sumd2t The sum of the distinct values read from the data file for the second temperature derivative of a given Pitzer theta coefficient. Dimensioning: sumd2t(npxpar). Usage: sumd2t(k) is the sum for the k-th pair of cations or pair of anions. This array is used

to compute the average values of the second temperature derivatives of the theta coefficients for such pairs (see **d2th**). This array is unique to EQPT.

- sumdt The sum of the distinct values read from the data file for the first temperature derivative of a given Pitzer theta coefficient. Dimensioning: sumdt(npxpar). Usage: sumdt(k) is the sum for the k-th pair of cations or pair of anions. This array is used to compute the average values of the first temperature derivatives of the theta coefficients for such pairs (see dth). This array is unique to EQPT.
- sumt The sum of the distinct values read from the data file for a given Pitzer theta coefficient. Dimensioning: sumt(npxpar). Usage: sumt(k) is the sum for the k-th pair of cations or pair of anions. This array is used to compute the average values of the theta coefficients for such pairs (see atheta). This array is unique to EQPT.
- summ The number of distinct values read from the data file for a given Pitzer theta coefficient. Dimensioning: summ(npxpar). Usage: summ(k) is the number of distinct values for the k-th pair of cations or pair of anions. This array is used to compute the average values of the theta coefficients and the average values of the first and second temperature derivatives for such pairs (see atheta, dth, and d2th). This array is unique to EQPT.
- tdamax The nominal upper temperature limit of the data file, °C.
- tdamin The nominal lower temperature limit of the data file, °C.
- **tempc** Temperature, °C, on the standard temperature grid. Dimensioning: **tempc(8)**. Usage: **tempc(n)** is the temperature for the **n**-th point on the grid. Note: dimensioning and usage differ from that in EQ3NR and EQ6.
- tempcs The scaled temperature on the standard temperature grid. Dimensioning: tempcs(8). Usage: tempcs(n) is the scaled temperature for the n-th point on the grid. This is used in fitting interpolating polynomials to data on the standard grid. The scaled temperature array is computed from the tempc array by normalizing the values on each segment of the grid by dividing by the maximum value on that segment. This array is unique to EQPT.
- ${}^{S}\theta_{MN}$ thdum The Pitzer short range theta coefficient (25°C). This is a local variable in module rdpz3.f used to read in the value of this coefficient for the current mixture of two aqueous neutral electrolytes containing a common ion. See theta and atheta. This variable is unique to EQPT.

 ${}^{S}\theta_{MN}$ theta The Pitzer short range theta coefficient (25°C). This is an array in which is saved the theta coefficient for each mixture of two aqueous neutral electrolytes containing a common ion. Dimensioning: theta(ixspar,npxpar). Usage: theta(i,nmx) is the i-th theta value read for the nmx-th cation-cation or anion-anion pair. This array is actually presently not used for anything. It could be used to summarize the situation when the theta values read for a given pair of cations or anions are not all identical. See thdum and atheta. This array is unique to EQPT.

- tmaxThe max norm of the temperature on a range of the standard temperature grid. See
tempsc.This variable is unique to EQPT.
- **ubdotp** Array of lines read from the data file which specify hard core diameters and **insgfl** (**ineu**) flags for the aqueous species.

ucode	A variable containing the name of the code.				
uelem	Array of names of chemical elements (their chemical symbols). Dimensioning: uelem(nctpar) . Usage: uelem(nc) is the name of the nc -th element.				
uendit	The string 'endit.'.				
ueqlrn	A string containing the release number of the supporting EQLIB library.				
ueqlst	A string containing the stage number of the supporting EQLIB library.				
uhead	Data file type key string: = 'stfipc' Simple extended Debye-Hückel formalism = 'stpitz' Pitzer formalism				
ukey	Data file key string: The value read from the data file is the data file key, which is one of the following: = 'com' Composite = 'sup' SUPCRT92 = 'nea' NEA = 'hmw' Harvie-Møller-Weare (Harvie, Møller, and Weare, 1984) = 'pit' Pitzer (Pitzer, 1979)				
	These are mapped to one of the following: = 'bdot' Simple extended Debye-Hückel formalism ('com', 'sup', "nea") = 'pitzer' Pitzer formalism ('hmw", 'pit')				
	Functionally, these latter values duplicate the function of the uhead key string.				
umsp	Array containing the names of the cations corresponding to pure aqueous neutral electrolytes read from the data file. The anions are stored in uxsp . Instead of a cation-anion pair, the code will also accept a pair of neutral species, in which case one will be stored in umsp , the other in uxsp . Dimensioning: umsp(npxpar) . Usage: umsp(npxp) is the cation (or one of two neutral species) in the npxp -th pure aqueous neutral electrolyte or pair of neutral species. This array is unique to EQPT.				
unam24	Array containing the names of the species appearing in a chemical reaction given on the data file. Dimensioning: unam24(nsqpa1) . Usage: unam24(n) is the name of the species corresponding to the n -th coefficient in the current reaction as it is written on the data file. This array is known as udrxd in EQ3NR and EQ6.				
unam8	Holding array of names of chemical elements. Dimensioning: unam8(nctpar) . Us- age: unam8(n) is the name of the n -th element listed for the current species. See cdum1 . This array is known as uelemd in EQ3NR and EQ6.				
unone	The string 'none'.				
uoxide	Array of names of oxides of the chemical elements. Dimensioning: uoxide(nctpar) . Usage: uoxide(nc) is the name of the oxide of the nc -th chemical element. This array is presently vestigial. See also oxide .				
urelno	A string containing the release number of the code it is contained in.				
ustage	A string containing the stage number of the code it is contained in.				

utitld	The title (text) from the data1 file. Dimensioning: utitld(ntitpa) . Usage: utitld(n) is the n -th line of this title.
uxsp	Array containing the names of the anions corresponding to pure aqueous neutral electrolytes read from the data file. The cations are stored in umsp . Instead of a cation-anion pair, the code will also accept a pair of neutral species, in which case one will be stored in umsp , the other in uxsp . Dimensioning: uxsp(npxpar) . Usage: uxsp(npxp) is the anion (or one of two neutral species) in the npxp -th pure aqueous neutral electrolyte or pair of neutral species. This array is unique to EQPT.
uref	Array of reference strings for the Pitzer theta coefficient for a pair of cations or a pair of anions. Dimensioning: uref(npxpar) . Usage: uref(nmx) is the reference string for the nmx -th such pair. This array is unique to EQPT. Note: uref is also used as a local variable with a slightly different meaning in module pcrss.f .(reference string for a solid solution).
urefp	Array of reference strings for the Pitzer coefficient data for mixtures of two aqueous neutral electrolytes. Dimensioning: uref(npxpar) . Usage: uref(npxp) is the reference string for the npxp -th such mixture.
usp1	Array of names of the first of a pair of ions of the same charge sign for which the data file contains a Pitzer theta coefficient. Dimensioning: usp1(npxpar) . Usage: ups1(nmx) is the name of the first species in the nmx -th such pair. See also usp2 . This array is unique to EQPT. Note: usp1 and usp2 are also used as local variables for species names in module pdpz2.f .
usp2	Array of names of the second of a pair of ions of the same charge sign for which the data file contains a Pitzer theta coefficient. Dimensioning: usp1(npxpar) . Usage: ups1(nmx) is the name of the second species in the nmx -th such pair. See also usp1 . This array is unique to EQPT. Note: usp1 and usp2 are also used as local variables for species names in module pdpz2.f .
uspcm	Array of names of the third of a triplet of ions composing two aqueous neutral elec- trolytes containing a common ion. This third ion has the opposite charge sign of the other two. Dimensioning: uspcm(ixspar,npxpar) . Usage: uspcm(ixs(nmx),nmx) is the name of the third species in the ixs(nmx) -th triplet containing the nmx -th dis- tinct pair of the first two ions. See also theta and ixs . This array is unique to EQPT.
uspec	Array of names of species. Dimensioning: uspec(nsqpa1) . Usage: uspec(ns) is the name of the ns- th species. The first nsq species are the aqueous basis species. The nsq1- th species is the current non-basis species. This may be a species belonging to any type of phase. Dimensioning and usage differ in EQPT from that in EQ3NR and EQ6.
uspn	Holding array which contains the names of two consecutively read species of a giv- en phase type. Dimensioning: uspn(2). Usage: uspn(1) and uspn(1) are the names of two consecutive such species. This array is used to write species lists to the out- put and slist files. It is unique to EQPT.
uspp1	Array of names of the first of a triplet of ions composing two aqueous neutral elec- trolytes containing a common ion. This first ion is one of two having the same charge sign. Dimensioning: uspp1(npxpar) . Usage: uspp1(nmpp) is the name of the first species in the nmpp -th such triplet. See also uspp2 and usppc . This array is unique to EQPT.

	uspp2	Array of names of the second of a triplet of ions composing two aqueous neutral electrolytes containing a common ion. This second ion is one of two having the same charge sign. Dimensioning: uspp2(npxpar) . Usage: uspp2(nmpp) is the name of the second species in the nmpp -th such triplet. See also uspp1 and usppc . This array is unique to EQPT.
	usppc	Array of names of the third of a triplet of ions composing two aqueous neutral elec- trolytes containing a common ion. This third ion has the opposite charge sign of the other two. Dimensioning: usppc(npxpar) . Usage: usppc(nmpp) is the name of the third species in the nmpp -th such triplet. See also uspp1 and uspp2 . This array is unique to EQPT.
	ussp	Array of names of the ions (or neutral species) defining an aqueous neutral electro- lyte (or pair of neutral species) for which Pitzer coefficient data are read from the data file. This array contains no duplications. Dimensioning: ussp(npxpar) . Usage: ussp(j) is the name of the j -th species in this array. This array is created by extracting the species names from the ussrs array. See also nssp . This array is presently not used for anything. It array is unique to EQPT.
	ussrs	Array of pairs of names of the ions (or neutral species) defining an aqueous neutral electrolyte (or pair of neutral species) for which Pitzer coefficient data are read from the data file. The first ion is normally the cation. Dimensioning: ussrs(2,npxpar) . Usage: ussrs(1,(npxp) is the name of the first species in the npxp -th such pair; ussrs(2,(npxp) is the name of the second species. This array is unique to EQPT.
	xbarlm	Array of limits on the mole fractions of solid solution end-member components. Di- mensioning: xbarlm(iktpar) . Usage: xbarlm(ik) is the limit on the mole fraction of the ik -th component of the current solid solution.
log K _{Eh}	xlkeh	Log equilibrium constant of the half reaction relating the hypothetical electron and $O_{2(g)}$. Dimensioning: xlkeh(8) . Usage: xlkeh(n) is the log equilibrium constant of this half reaction for the n -th point on the standard temperature grid. Note: dimensioning and usage differ from that in EQ3NR and EQ6.
log K _r	xlks	Array of log equilibrium constants of the dissociation/destruction reactions of aque- ous species. Dimensioning: xlks(8,nsqpa1) . Usage: xlks(n,ns) is the log of the equi- librium constant for the n -th point on the standard temperature grid for the ns -th species. The first nsq species are the aqueous basis species. The nsq1 -th species is the current non-basis species and may be of any phase type. Note: dimensioning and usage differ from that in EQ3NR and EQ6.
z _i	Z	Array of electrical charges of the species on the data file. Dimensioning: z (nsqpa1). Usage: z (ns) is the electrical charge of the ns -th species. The first nsq species are the aqueous basis species. The nsq1 -th species is the current non-basis species and may be of any phase type. Dimensioning and usage differ in EQPT from that in EQ3NR and EQ6.
å _i	zero	Hard core diameter of the current aqueous species. See ubdotp . Note: the array azero is used to hold hard core diameters in EQ3NR and EQ6. Here ' zero ' appears to be a corruption of azero .
	zm	Array of electrical charges of the first species (cation or neutral) in an aqueous neu- tral electrolyte (or pair of neutral species) for which Pitzer coefficient data are read from the data file. Dimensioning: zm (npxpar). Usage: zm (npxpar) is the electrical

charge of the first species in the **npxp**-th such aqueous neutral electrolyte or pair of neutral species. See also **zx**.

Array of electrical charges of the second species (anion or neutral) in an aqueous neutral electrolyte (or pair of neutral species) for which Pitzer coefficient data are read from the data file. Dimensioning: zx(npxpar). Usage: zx(npxpar) is the electrical charge of the second species in the npxp-th such aqueous neutral electrolyte or pair of neutral species. See also zm.

Appendix B. Glossary of EQPT Subroutines

EQPT is a medium-sized code. The source code consists of the main program and a number of subroutines. In addition, EQPT uses several modules from the EQLIB library. These are described in Appendix B of the EQ3/6 Package Overview and Installation Guide (Wolery, 1992a). The modules are described as ".f" files, as this is how they are normally worked with under a UNIX operating system. For a description of the code architecture, see Chapter 5.

bldsp.f	This module is called by the EQPT module eqpt.f . It builds the array ussp from the ussrs array. The array ussp contains the names of all species which appear in connection with Pitzer coefficient data for pure aqueous neutral electrolytes (cation-anion pair) or pairs of neutral species. Names that are found more than once in ussrs are copied only once to ussp . The ussp array is presently not used for anything.
eqpt.f	This module is the main program of EQPT. It calls other modules to read the data file, process the data, and write the data1 and other output files.
funcy.f	This module is called by the EQPT module intrp.f . It evaluates fitted interpolating polynomials at the points corresponding to the standard temperature grid. It expects one interpolating polynomial for the range 0-100°C, another for the range 100-300°C.
gnenb.f	This module is called by the EQPT module eqpt.f . It scans the data file, counting the number of chem- ical elements (nct) and the number of basis species (nsq). It the rewinds the data file.
gridpy.f	This module is called by the EQPT modules pcraq.f and pcrsg.f . It writes specified data on the stan- dard temperature grid to a specified file.
intrp.f	This module is called by the EQPT modules pcraq.f , pcrsg.f , and wrpar.f . It fits interpolating polynomials to specified data on the standard temperature grid. It fits one interpolating polynomial for the range 0-100°C, another for the range 100-300°C.
ofiles.f	This module is called by the EQPT module eqpt.f . It opens all files, except for output , which is opened by eqpt.f .
pcraq.f	This module is called by the EQPT module eqpt.f . It reads the composition, reaction, and standard state thermodynamic data for all aqueous species. It checks the data for simple errors, such as unbalanced reactions.
pcrsg.f	This module is called by the EQPT module eqpt.f . It reads the composition, reaction, and standard state thermodynamic data for all pure mineral and gas species. It checks the data for simple errors, such as unbalanced reactions.
pcrss.f	This module is called by the EQPT module eqpt.f. It reads the data for all solid solutions.
pdpz2.f	This module is called by the EQPT module eqpt.f . It reads the data for all Pitzer coefficient parameters associated with pure aqueous neutral electrolytes or pairs of aqueous neutral species. It does limited processing of these data. It writes these data on the dpt1 file.
preacy.f	This module is called by the EQPT modules pcraq.f , pcrsg.f , and rxnchk.f . It writes a specified chemical reaction onto a specified file.

rdpar.f	This module applicable:	is called by the EQPT module eqpt.f. It reads the data for the following data grids, as			
	tempc press	standard temperature grid pressure on the standard temperature grid			
	_				
	adh bdh	Debye-Hückel A_{γ} parameter on the standard temperature grid Debye-Hückel B_{γ} parameter on the standard temperature grid			
	bdot cco2	extended Debye-Hückel \dot{B} parameter on the standard temperature grid coefficients of the Drummond (1981) polynomial for the log activity coefficient of dissolved CO_2			
	aphi	Debye-Hückel A_{ϕ} parameter on the standard temperature grid			
	xlkeh	$\log K_{Eh}$ on the standard temperature grid			
rdpz3.f		is called by the EQPT module eqpt.f . It reads the data for all Pitzer coefficient parameters ith mixtures of two aqueous neutral electrolytes It does limited processing of these data.			
rdwele.f	This module is called by the EQPT module eqpt.f . It reads the data for the chemical elements and writes it on the data1 file.				
rdwttl.f	This module is called by the EQPT module eqpt.f . It reads the title of the data file and writes it on the data1 file.				
rdbdot.f	This module is called by the EQPT module eqpt.f . It reads the lines of the data file which contain hard core diameters and insgfl flags for the aqueous species. These data are used in conjunction with the B-dot equation for the activity coefficients of aqueous species.				
rxnchk.f	This module is called by the EQPT modules pcraq.f and pcrsg.f . It checks a specified reaction for mass and charge balance. If an error is found, an error message is written to the output and screen files.				
srch22.f	This module is called by the EQPT module wrpz3.f . It searches to parallel arrays of species names to see if a common index exists which corresponds to an input pair of species names. If such an index exists, it is returned. It also returns information on which array each input species name was found in.				
wrbdot.f	This module is called by the EQPT module eqpt.f . It writes the data for the hard core diameters and insgfl flags for the aqueous species onto the data1 file. The lines containing these data are read from the data file by module rdbdot.f .				
wrpar.f		is called by the EQPT module eqpt.f . It processes the data read from the data file by mod- It then writes the corresponding data onto the data1 file.			
wrpz3.f		is called by the EQPT module eqpt.f . It completes the processing of all Pitzer coefficient tes the results onto the data1 file.			

Appendix C. EQPT Error Messages

All EQ3/6 error messages fit into one of three categories: *error*, *warning*, and *note*. An *error* implies a fatal error. Execution of the current problem will cease without completion, immediately in some cases, later in others. Which is the case depends on whether it makes more sense to stop immediately or to continue checking for other errors before ceasing execution. A *warning* indicates a condition which may or may not represent a real error. A *note* indicates a condition knowledge of which may assist the user in determining or tracking down the cause of an error. An *error* message may be preceded or followed by one or more other *error*, *warning*, or *note* messages. These messages are written to both the screen file and the **output** file.

Each EQ3/6 error message has the following format:

* msgtype - (source/module) Message.

where msgtype = error, warning, or *note*, *source* is the root name of the source file (e.g., *eqlib*, *eqpt*, *eq3nr*, or *eq6*) containing the *module*, *module* is the name of the module (main program or subroutine) which writes the message, and *Message* is the message itself. The messages are designed to be as self-explanatory as possible. The messages are reproduced here using AAAA to stand for a character variable, *IIII* for an integer, and *RRRR* for a floating point number.

EQPT has no input file. Most of the error messages that users are likely to encounter pertain to problems regarding the **data0** data file that this code is processing. In most instances, the meaning of these messages should be immediately clear to the user. In other instances, it may be necessary to search out other information. In such cases, there are three principal actions that users should take. The first is to check the **output** file for additional diagnostic messages (*warnings* and *notes*) which may bear on the matter. You may find that certain data in the **data0** file were not entered in the correct fields, or that a line is missing or you have an extra line. User's should expect to see such errors only if they locally modify the data files supplied as part of the EQ3/6 export package or if they make up their own data files. Otherwise, the occurrence of an error is probably due to corruption of the files in the export package.

Some messages deal with programming errors. The user should see these rarely if ever. These are likely to appear somewhat more cryptic to users. Problems of this type must be dealt with by diagnosing the problem (probably with the help of a symbolic debugger) and modifying the code. Most users should probably not attempt corrective action of this sort. The code custodian should be notified of suspected programming errors and may be able to provide fixes.

Some of the messages displayed in this appendix are followed by *Comments* that may help to explain them. The list of messages given here include only those generated by EQPT modules. Users of EQPT may also encounter error messages from EQLIB modules. These messages are listed in similar format in Appendix C of the EQ3/6 Package Overview and Installation Guide (Wolery, 1992a). The *errors* are listed first, then the *warnings* and finally the *notes*.

Message: * error - (eqpt/eqpt) Unrecognized data file key = "AAAA". Allowed values are "bdot", "hmw", and "pitzer".

- Comment: In order to read the data file correctly, the code must know what kind of activity coefficient data to read. The two present data file archetypes are represented by the **com** and **hmw** data files. The archetype is presently defined by the **ukey** string. The normal data file key string (**com**, **sup**, **nea**, **hmw**, or **pit**) is mapped in module **rdwttl.f** to **bdot** for data files of the **com** archetype, and to **pitzer** for files of the **hmw** archetype. The **ukey** value of **hmw** is not actually bused, but is equivalent to **pitzer**. Check the data file to see that the normal data file key string is not missing. If you have created your own such key string, you must declare it in module **rdwttl.f**.
- Message: * error (eqpt/gnenb) Number of chemical elements on the data file exceeds the dimensioned limit (nctpar) of IIII.
- Comment: Increase the dimensioned limit.
- Message: * error (eqpt/gnenb) End-of-file hit or other read error occurred while reading the AAAA.
- Comment: Search the **output** file for previous error, warning, or note messages. Check the end of the **data0** file to see that it has not been abnormally truncated. Also check the **data0s** file (the copy of **data0** that has been stripped of comments); this is what the code was actually reading.
- Message: * error (eqpt/gnenb) Number of expanded basis species exceeds the dimensioned limit (nsqpar) of IIII.
- Comment: Increase the dimensioned limit. The expanded basis set is the total basis set, including the strict and auxiliary basis sets.
- Message: * error (eqpt/intrp) Could not compute the coefficients of an interpolating polynomial in the temperature range 0-100 Celsius.
- Comment: Check the data grid that was being processed.
- Message: * error (eqpt/intrp) Could not compute the coefficients of an interpolating polynomial in the temperature range 100-300 Celsius.
- Comment: Check the data grid that was being processed.
- Message: * error (eqpt/pcraq) Species "AAAA" is composed of no chemical elements.
- Comment: Check the elemental composition of the species specified on the data file. If it is okay, look for a missing or extra line immediately above it.
- Message: * error (eqpt/pcraq) Species "AAAA" is composed of IIII chemical elements, but there are only IIII elements on the data file.
- Comment: This is most likely due to a typographical error in specifying the composition of the species. Check the specified composition. If it seems okay, check it against the chemical elements block. If that seems okay, look for a missing or extra line immediately above the composition data for the species for which the problem occurred.
- **Message:** * error (eqpt/pcraq) Unrecognized chemical element "AAAA" is listed in the composition of species "AAAA".
- Comment: The data file may not contain a species composed of an element that has not been declared in the chemical elements block. This message is most likely to be encountered, however, because of a typographical error in the specification of the elemental composition of the species (e.g., "ci" or "cu" for "cu" when specifying a stoichiometric coefficient for copper).

- Message: * error (eqpt/pcraq) The reaction for the destruction of species "AAAA" includes IIII species, but there are only IIII basis species on the data file, so only IIII species may appear in the reaction.
- Comment: This is most likely due to a typographical error in specifying the associated reaction of the species. Check the specified reaction. If it seems okay, check it against the set of basis species. If that seems okay, look for a missing or extra line immediately above the reaction for the species for which the problem occurred.
- Message: * error (eqpt/pcraq) Non-basis species "AAAA" has no species in the reaction in which it is destroyed.
- Comment: This is most likely due to a typographical error in specifying the associated reaction of the species. Check the specified reaction. If it seems okay, look for a missing or extra line immediately above the reaction.
- **Message:** * error (eqpt/pcraq) The reaction which destroys non-basis species "AAAA" is written in terms of an unrecognized basis species "AAAA".
- Comment: This is most likely due to a typographical error in specifying the associated reaction of the species (e.g., "ci++" or "cu++" for "cu++"). Check the specified reaction. If it seems okay, check it against the set of basis species. If that seems okay, look for a missing or extra line immediately above the reaction for the species for which the problem occurred.
- Message: * error (eqpt/pcrsg) Species "AAAA" is composed of no chemical elements.
- Comment: Check the elemental composition of the species specified on the data file. If it is okay, look for a missing or extra line immediately above it.
- Message: * error (eqpt/pcrsg) Species "AAAA" is composed of IIII chemical elements, but there are only IIII elements on the data file.
- Comment: This is most likely due to a typographical error in specifying the composition of the species. Check the specified composition. If it seems okay, check it against the chemical elements block. If that seems okay, look for a missing or extra line immediately above the composition data for the species for which the problem occurred.
- **Message:** * error (eqpt/pcrsg) Unrecognized chemical element "AAAA" is listed in the composition of species "AAAA".
- Comment: The data file may not contain a species composed of an element that has not been declared in the chemical elements block. This message is most likely to be encountered, however, because of a typographical error in the specification of the elemental composition of the species (e.g., "ci" or "cu" for "cu" when specifying a stoichiometric coefficient for copper).
- **Message:** * error (eqpt/pcrsg) The reaction for the destruction of species "AAAA" includes IIII species, but there are only IIII basis species on the data file, so only IIII species may appear in the reaction.
- Comment: This is most likely due to a typographical error in specifying the associated reaction of the species. Check the specified reaction. If it seems okay, check it against the set of basis species. If that seems okay, look for a missing or extra line immediately above the reaction for the species for which the problem occurred.
- Message: * error (eqpt/pcrsg) Non-basis species "AAAA" has no species in the reaction in which it is destroyed.
- Comment: This is most likely due to a typographical error in specifying the associated reaction of the species. Check the specified reaction. If it seems okay, look for a missing or extra line immediately above the reaction.

- **Message:** * error (eqpt/pcrsg) The reaction which destroys non-basis species "AAAA" is written in terms of an unrecognized basis species "AAAA".
- Comment: This is most likely due to a typographical error in specifying the associated reaction of the species (e.g., "ci++" or "cu++" for "cu++"). Check the specified reaction. If it seems okay, check it against the set of basis species. If that seems okay, look for a missing or extra line immediately above the reaction for the species for which the problem occurred.
- Message: * error (eqpt/pcrss) Solid solution "AAAA" is composed of IIII end-members, which exceeds the maximum dimension (iktpar) of IIII.
- Comment: Increase the dimensioned limit.
- Message: * error (eqpt/pdpz2) The number of single salt entries exceeds the dimension limit (npxpar) of IIII.
- Comment: Increase the dimensioned limit. A "single salt entry" refers to a block of Pitzer coefficient data for a pure aqueous neutral electrolyte or a pair of aqueous neutral species.
- Message: * error (eqpt/pdpz2) End-of-file hit or other read error occurred while reading single salt parameters.
- Comment: The "single salt parameters" refer to the superblock of Pitzer coefficient data for pure aqueous neutral electrolytes or pairs of aqueous neutral species. Search the **output** file for previous error, warning, or note messages. Check the end of the **data0** file to see that it has not been abnormally truncated. Also check the **data0s** file (the copy of **data0** that has been stripped of comments); this is what the code was actually reading.
- Message: * error (eqpt/rdbdot) The number of bdot lines on the data file exceeds the dimensioned limit (nstpar) of IIII.
- Comment: Increase the dimensioned limit. The "bdot lines" are the lines in the block of activity coefficient data for data files of the **com** archetype. They contain the hard core diameters and neutral species treatment flags for the aqueous species on the data file.
- Message: * error (eqpt/rdbdot) End-of-file hit or other read error occurred while reading bdot parameters.
- Comment: The "bdot lines" are the lines in the block of activity coefficient data for data files of the **com** archetype. They contain the hard core diameters and neutral species treatment flags for the aqueous species on the data file. Search the **output** file for previous error, warning, or note messages. Check the end of the **data0** file to see that it has not been abnormally truncated. Also check the **data0s** file (the copy of **data0** that has been stripped of comments); this is what the code was actually reading.
- Message: * error (eqpt/rdpar) End-of-file hit or other read error occurred while reading data0 parameters.
- Comment: The "data0 parameters" refers to the blocks containing the temperatures on the standard grid, the pressure grid, the grids for the Debye-Hückel and related parameters, and the grid for the log equilibrium constant for the "Eh reaction.". Search the **output** file for previous error, warning, or note messages. Check the end of the **data0** file to see that it has not been abnormally truncated. Also check the **data0s** file (the copy of **data0** that has been stripped of comments); this is what the code was actually reading.
- **Message:** * error (eqpt/rdpz3) The species triplet of "AAAA", "AAAA", and "AAAA" representing two electrolytes with a common ion has one species appearing twice.
- Comment: This refers to an error in one of the blocks of Pitzer coefficient data for mixtures of two aqueous neutral electrolytes. The species triplet for one of these mixtures is improperly specified.

Message: * error - (eqpt/rdpz3) Psi array index overflow. The dimensioning parameter npxpar is too small.

Comment: Increase the dimensioned limit.

Message: * error - (eqpt/rdpz3) Theta array index overflow. The dimensioning parameter npxpar is too small.

Comment: Increase the dimensioned limit.

Message: * error - (eqpt/rdpz3) Have ixs array index overflow. The dimensioning parameter ixspar is too small.

Comment: Increase the dimensioned limit.

Message: * error - (eqpt/rdpz3) End-of-file hit or other read error occurred while reading mixture term parameters.

Comment: The "mixture term parameters" refer to the superblock of Pitzer coefficient data for mixtures of two aqueous neutral electrolytes. Search the **output** file for previous error, warning, or note messages. Check the end of the **data0** file to see that it has not been abnormally truncated. Also check the **data0s** file (the copy of **data0** that has been stripped of comments); this is what the code was actually reading.

Message: * error - (eqpt/rdwele) End-of-file hit or other read error occurred while reading chemical elements block.

- Comment: Search the **output** file for previous error, warning, or note messages. Check the end of the **data0** file to see that it has not been abnormally truncated. Also check the **data0s** file (the copy of **data0** that has been stripped of comments); this is what the code was actually reading.
- Message: * error (eqpt/rdwttl) Have read invalid data0 file header: AAAA

Comment: Check the header. It must begin with "data0".

- Message: * error (eqpt/rdwttl) End-of-file hit or other read error occurred while the data0 file title.
- Comment: Search the **output** file for previous error, warning, or note messages. Check the end of the **data0** file to see that it has not been abnormally truncated. Also check the **data0s** file (the copy of **data0** that has been stripped of comments); this is what the code was actually reading.
- Message: * error (eqpt/rxnchk) The following reaction has a computed electrical imbalance of RRRR-
- Comment: Check the reaction. If it seems okay, check the electrical charges specified for the species which appear in the reaction.
- Message: * error (eqpt/rxnchk) The following reaction has a computed mass imbalance for "AAAA" of RRRR-
- Comment: Check the reaction. If it seems okay, check the elemental compositions specified for the species which appear in the reaction.
- Message: * error (eqpt/wrpz3) Can not find the species pair "AAAA" and "AAAA" in the parallel species name arrays umsp and uxsp. Unable to process psi data.
- Comment: This is probably due to the specification of Pitzer coefficient data for a mixture of two aqueous neutral electrolytes, but without specification of the corresponding data for one or both electrolytes. If these data seem to be present, look for a typographical error in a species name (e.g., "na++" or "na+" for "na+").

- Message: * note (eqpt/pcrsg) The pure liquids block has not been written on the data1 and data1f files, because the EQ3NR and EQ6 codes presently do not treat non-aqueous liquids.
- Comment: In the future, EQ3/6 may be expanded to handle non-aqueous liquids. In the meantime, EQ3NR and EQ6 have no way to deal with thermodynamic data for such species/phases.

Message: * note - (wrpz3) alphas=99. since lambda1=lambda2=zero

Comment: This message is only written to the dpt1 file. It is written as part of the header for the Pitzer theta coefficients block. It is intended merely to note that the lambda coefficients obtained from the theta coefficients have no ionic strength dependence. This message is largely vestigial.

Appendix D. Known Bugs and Such

This appendix presents notes on known bugs and other known unusual phenomena.

1. EQPT does not currently provide for treating all types of observable interaction coefficients belonging to Pitzer's equations involving interactions with electrically neutral species. See Chapter 3 of the EQ3NR Theoretical Manual and User's Guide (Wolery, 1992b) for a discussion of the various types of observable interaction coefficients belonging to Pitzer's equations.

For a complete list of known bugs and such for EQ3/6, see Appendix D of the EQ3/6 Package Overview and Installation Guide (Wolery, 1992a).

Appendix F. The slist Files for the com and hmw Data Files

This appendix presents the **slist** ("species list") files for the **data0.com.R10** and **data0.hmw.R10** data files.

The <u>slist</u> file for the <u>data0.com.R10</u> data file:

EQPT Species List File:

no. of elements on the data file = 78
 the dimensioned limit = 100
no. of aqueous species in the master set = 147
 the dimensioned limit = 500

data0.com.R10 THERMODYNAMIC DATABASE generated by gembochs/INGRES 15-apr-91

- ·						1 - 0 0 0 4 0
element	=	0	,	atwt	=	15.99940
element	=	ag	,	atwt	=	107.86820
element	=	al	,	atwt	=	26.98154
element	=	am	,	atwt	=	243.00000
element	=	ar		atwt	=	39.94800
			'			
element	=	au	'	atwt	=	196.96654
element	=	b	,	atwt	=	10.81100
element	=	ba	,	atwt	=	137.32700
element	=	be	,	atwt	=	9.01218
element	=	br	,	atwt	=	79.90400
element	=	ca		atwt	=	40.07800
	Ξ		'			112.41100
element		cd	'	atwt	=	
element	=	ce	,	atwt	=	140.11500
element	=	cl	,	atwt	=	35.45270
element	=	CO	,	atwt	=	58.93320
element	=	cr	,	atwt	=	51.99610
element	=	CS		atwt	=	132.90543
element			'	atwt		63.54600
	=	cu	'		=	
element	=	dy	,	atwt	=	162.50000
element	=	er	,	atwt	=	167.26000
element	=	eu	,	atwt	=	151.96500
element	=	f	,	atwt	=	18.99840
element	=	fe		atwt	=	55.84700
			'			
element	=	ga	'	atwt	=	69.72300
element	=	gd	'	atwt	=	157.25000
element	=	h	,	atwt	=	1.00794
element	=	as	,	atwt	=	74.92159
element	=	С	,	atwt	=	12.01100
element	=	he		atwt	=	4.00206
element	=	hg	'	atwt	=	200.59000
-			'			
element	=	ho	'	atwt	=	164.93032
element	=	р	,	atwt	=	30.97362
element	=	i	,	atwt	=	126.90447
element	=	in	,	atwt	=	114.82000
element	=	k	,	atwt	=	39.09830
element	=	kr		atwt	=	83.80000
	_		'		-	138.90550
element		la	'	atwt		
element	=	li	,	atwt	=	6.94100
element	=	lu	,	atwt	=	174.96700
element	=	mg	,	atwt	=	24.30500
element	=	mn	,	atwt	=	54.93085
element	=	mo		atwt	=	95.94000
element	=	na	'	atwt	=	22.98977
			'			
element	=	nd	'	atwt	=	144.24000
element	=	ne	,	atwt	=	20.17970
element	=	ni	,	atwt	=	58.69000
element	=	n	,	atwt	=	14.00674
element	=	np	,	atwt	=	237.04800
element	=	pb		atwt	=	207.20000
element	=		'	atwt	=	106.42000
		pd	'			
element	=	pr	,	atwt	=	140.90765
element	=	pu	,	atwt	=	244.00000
element	=	ra	,	atwt	=	226.02500
element	=	rb	,	atwt	=	85.46780
element	=	re	,	atwt	=	186.20700
element	=	rn		atwt	=	222.00000
			'			101.07000
element	=	ru	'	atwt	=	
element	=	SC	,	atwt	=	44.95591
element	=	se	,	atwt	=	78.96000
element	=	si	,	atwt	=	28.08550
element				atwt	=	
	=	sm	,	alwi	_	150.36000
element			'			
element element	=	sn s	;	atwt atwt	=	118.71000 32.06600

element	=	sr	,	atwt	=	87.62000
element	=	tb	,	atwt	=	158.92534
element	=	tc	,	atwt	=	98.00000
element	=	th	,	atwt	=	232.03810
element	=	ti	,	atwt	=	47.88000
element	=	tl	,	atwt	=	204.38330
element	=	tm	,	atwt	=	168.93421
element	=	u	,	atwt	=	238.02890
element	=	v	,	atwt	=	50.94150
element	=	W	,	atwt	=	183.85000
element	=	xe	,	atwt	=	131.29000
element	=	У	,	atwt	=	88.90585
element	=	yb	,	atwt	=	173.04000
element	=	zn	,	atwt	=	65.39000
element	=	zr	,	atwt	=	91.22400

aqueous

queou	S	
1	h2o	201
3	al+++	ag+
		am+++
5 7	ar(aq)	au+
	b(oh)3(aq)	ba++
9	be++	br-
11	ca++	cd++
13	ce+++	cl-
15	CO++	cro4
17	cs+	cu++
19	dy+++	er+++
21	eu+++	f -
23	fe++	ga+++
25	gd+++	h+
27	h2aso4-	hco3-
29	he(aq)	hg++
31	ho+++	hpo4
33	i-	in+++
35	k+	kr(aq)
37	la+++	li+
39	lu+++	mg++
41	mn++	moo4
43	na+	nd+++
45	ne(aq)	ni++
47	no3-	np++++
49	pb++	pd++
51	pr+++	pu++++
53	ra++	rb+
55	reo4-	rn(aq)
57	ruo4	sc+++
59	seo3	sio2(aq)
61	sm+++	sn++
63	so4	sr++
65	tb+++	tco4-
67	th++++	ti(oh)4(aq)
69	tl+	tm+++
71	uo2++	vo++
73	wo4	xe(aq)
75	у+++	yb+++
77	zn++	zr(oh)2++
79	o2(g)	(o-phth)
81		
83	acetic acid(aq)	acetone(aq)
85	ag++	am++++ amo2++
87	amo2+	
87	au+++	benzene(aq)
89 91	clo4-	CO+++
	co2(aq)	co3
93	cr++	cr+++
95	cro4	cu+
97	ethane(aq)	eu++
99	fe+++	glycine(aq)
101	h2(aq)	h2aso3-
103	h2po4-	hg2++
105	hs-	io3-
107	methanamine(aq)	methane(aq)
109	methanol(aq)	mn+++
111	mno4	n2(aq)
113	n3-	nh4+
115	no2-	np+++
117	npo2+	npo2++
119	o2(aq)	oh-
121	pb++++	po4
123	pu+++	puo2+
125	puo2++	ru(oh)2++
127	ru++	ru+++
129	ruo4(aq)	ruo4-
131	se	seo4
133	sm++	sn++++
135	so3	tc+++
137	tco++	tco4
139	tco4	u+++
141	u++++	uo2+
143	V+++	vo2+

145 147 149 151 153 155 157 159 161 163 165 167 171 173 175 177 179 181 183 185 187 191 193 195 197 201 203 205 227 209 221 223 225 227 229 221 223 225 227 229 221 223 235 227 229 221 223 225 227 229 221 223 235 227 229 221 223 225 227 229 221 223 225 227 229 221 223 235 227 229 221 223 225 227 229 221 223 235 227 229 221 223 225 227 229 221 223 235 227 229 221 223 225 227 229 221 223 225 227 229 221 223 225 227 229 221 223 225 227 229 221 223 225 227 229 221 223 225 227 229 221 223 225 227 229 221 223 225 227 229 221 223 225 227 229 221 223 235 227 229 221 223 235 227 229 221 223 235 227 229 221 223 225 227 229 221 223 225 227 229 221 223 225 227 229 221 223 235 227 229 221 223 225 227 229 221 223 225 227 229 221 223 225 227 229 221 223 225 227 229 221 223 235 227 229 221 223 225 227 229 221 223 225 227 229 221 223 225 227 229 221 223 225 227 229 221 223 225 227 229 221 223 225 227 229 221 223 225 227 229 221 223 225 227 229 221 223 225 227 229 221 223 225 227 229 227 229 221 223 225 227 229 227 227	<pre>vo4 zr++++ (npo2)3(oh)5+ (uo2)3(oh)5+ (uo2)3(co3)6(oh)12 (uo2)3(co3)6(6-) (uo2)3(oh)7- (vo)2(oh)2++ 1-butanol(aq) 1-butyne(aq) 1-betyne(aq) 1-betyne(aq) 1-betyne(aq) 1-betyne(aq) 1-pentyne(aq) 1-pentyne(aq) 1-pentyne(aq) 2-betyne(aq) 2-betyne(aq) 2-cotanone(aq) a-aminobutyric(aq) ag(co3)2 agc12- agc14 agf(aq) al(o-phth)+ al(oh)3(aq) alf*+ alanine(aq) alf*+ alanine(aq) alf*+ alanine(aq) alf*+ aminob++ am(co3)2- am(b)2+ ami(b)2+ ami(b)</pre>	yb- (ng (tc (uc (uc (uc (uc (uc 1-H 1-H 1-H 1-H 1-H 1-F 2-H 2-F 2-F 2-F 2-F 2-F 3-C 2-F 3-C 2-F 3-C 2-F 1-F 1-F 1-F 1-F 1-F 1-F 1-F 1-F 1-F 2-H 2-F 2-F 3-C 2-F 3-C 2-F 3-C 2-F 3-C 2-F 3-C 2-F 3-C 2-F 3-C 2-F 3-C 3-C 2-F 3-C 3-C 3-C 3-C 3-C 3-C 3-C 3-C 3-C 3-C
201	agf(aq)	agr
203	al(o-phth)+	al
205	al(oh)3(aq)	al
209	al2(oh)2++++	ali
211	alanine(aq)	alc
213	alf++	alf
215	alf3(aq)	alf
219	aloh++	als
221	am(co3)2-	am
223	am(h2po4)2+	am
229	am(so4)2-	amo
231	amcl2+	amo
233	amf++	amf
237 239 241 243	amno3++ amoh++ as(oh)3(aq) aso2-	ama ama asi
247	asparagine(aq)	ası
249	ass2-	b2c
251	b3o3(oh)4-	b4c
253	ba(o-phth)(aq)	bak
255 257 259 261 263	bach3coo+ baco3(aq) bano3+ beo2 bf3oh-	bad bad bad bf2
265	bh4-	bo2
267	br3-	bro
269	bro3-	bro
271	butanoate	but
273	ca(h3sio4)2(aq)	ca
275	cab(oh)4+	cac
277	cacl+	cac
279	caco3(aq)	cai
281	cah2po4+	cai
283	cah3sio4+	cal
285	cahp2o7-	cal
287	cano3+	cac
289	cap2o7	car
291	caso4(aq)	car
293	clo-	clo
295	clo3-	cn-
297	co(hs)2(aq)	co
299	co(oh)4	co2
301	co4(oh)4++++	co2
303	cocl+	col
305	coi2(aq)	cor
307	cos2o3(aq)	cos
309	coso4(aq)	cr
311 313 315 317 319	cr(oh)3(aq) cr2(oh)2++++ cr3(oh)4(5+) crcl++ cro3cl-	cri cri cri cri
321	csbr(aq)	cs
323	csi(aq)	cu
325	cu(nh3)2+	cu
327	cu(nh3)3++	cu
329	cucl+	cuo

++ >++
hpo2)2(oh)2++
puo2)2(oh)2++
tco(oh)2)2(aq)
uo2)2(oh)2++ 102)2(01)2++ 102)3(0h)4++ 102)3(0h)5co2+ 102)4(0h)7+ -butanamine(aq) butene(aq) heptanamine(aq) heptene(aq) hexanamine(aq) -hexene(aq) -octanamine(aq) octene(aq) pentanamine(aq) pentene(aq) propanamine(aq) propene(aq) butanone(aq) hexanone(aq) pentanone(aq) etate gcl(aq) gcl3--gco3no3(aq) (oh)2+ (oh)4- $(301)^{4}$ $(304)^{4}(301)^{24}(7+)$ $(301)^{4}(5+)$ $(301)^{4}(5+)$ $(301)^{4}(5+)$ $(301)^{4}(5+)$ f2+ f4hpo4+ .so4+ 1(co3)3---(h2po4)3(aq) 1(no3)2+ 1(oh)3(aq) 1(01) 1cl++ 1co3+ 1f2+ h2po4++ h02hco3(aq) nso4+ sh3(aq) o2oh-spartic acid(aq) so(oh)5-so5(oh)4-b(oh)4+ acl+ af+ aoh+ 2(oh)2-54-52-0--64 ro4-itanoic acid(aq) a(o-phth)(aq) ach3coo+ acl2(aq) f+ h2sio4(aq) ahco3+ ahpo4(aq) oh+ lpo4-lseo4(aq) .02o(oh)2(aq) 2(oh)3+ br2(aq) hs+ ons+ ono3+ oseo4(aq) c(oh)2+ c(oh)4-207--br++ cl2+ oh++ cl(aq) (co3)2--(nh3)2++ (no2)2(aq) cl2(aq)

331	1.0	1.2
222	cucl2-	cucl3
333	cucl4	cuco3(aq)
335 337	cuco3(oh)2	cuf+
339	cuh2po4+ cunh3+	cuhpo4(aq) cunh3++
341	cuno2+	cuo2
343	cuoh+	cupo4-
345	ethanamine(aq)	ethanol(aq)
347	ethylbenzene(aq)	ethylene(aq)
349	ethyne(aq)	eu(co3)2-
351	eu(co3)3	eu(hco3)++
353	eu(oh)(co3)2	eu(oh)2+
355	eu(oh)2co3-	eu(oh)3(aq)
357	eu(oh)4-	eu(oh)co3(aq)
359	eu(so4)2-	eu2(oh)2++++
361	eubr++	eubr2+
363	eubro3++	eucl++
365	eucl2+	euco3+
367	euf++	euf2+
	euio3++	euno3++
371	euoh++	euso4+
373	fe(ch3coo)2(aq)	fe(ch3coo)2+
375	fe(ch3coo)3(aq)	fe(oh)2(aq)
377	fe(oh)2+	fe(oh)3(aq)
379	fe(oh)3-	fe(oh)4-
381	fe(so4)2-	fe2(oh)2++++
383	fe3(oh)4(5+)	fech3coo+
385	fech3coo++	fecl+
387	fecl++	fecl2(aq)
389 391	fecl2+ fecl4	fecl4-
393	feco3+	feco3(aq) fef+
395	fef++	fef2+
397	feh2po4+	feh2po4++
399	fehco3+	fehpo4(aq)
401	fehpo4+	fehseo3++
403	fehso4++	feno2++
405	feno3++	feoh++
407	fepo4-	feso4(aq)
409	feso4+	formate
411	formic acid(aq)	glutamic acid(aq)
413	glutamine(aq)	h(o-phth)-
415	h2(o-phth)(aq)	h2cro4(aq)
417	h2f2(aq)	h2p2o7
419	h2po3f(aq)	h2s(aq)
421	h2se(aq)	h2seo3(aq)
423	h2sio4	h2so3(aq)
425	h2so4(aq)	h2vo4-
427	h3aso3(aq)	h3aso4(aq)
429	h3p2o7-	h3po4(aq)
431	h4(h2sio4)4	h4p2o7(aq)
433	h6(h2sio4)4	haso2(aq) haso4
435	haso3f-	
435 437	hass2(aq)	hbro(aq)
435 437 439	hass2(aq) hcl(aq)	hbro(aq) hclo(aq)
435 437 439 441	hass2(aq) hcl(aq) hclo2(aq)	hbro(aq) hclo(aq) hcoo2-
435 437 439 441 443	hass2(aq) hcl(aq) hclo2(aq) hcro4-	hbro(aq) hclo(aq) hcoo2- heptanoate
435 437 439 441 443 445	hass2(aq) hcl(aq) hclo2(aq) hcro4- heptanoic acid(aq)	hbro(aq) hclo(aq) hcoo2- heptanoate hexanoate
435 437 439 441 443 445	hass2(aq) hcl(aq) hclo2(aq) hcro4- heptanoic acid(aq) hexanoic acid(aq)	hbro(aq) hclo(aq) hcoo2- heptanoate hexanoate hf(aq)
435 437 439 441 443 445 447	hass2(aq) hcl(aq) hclo2(aq) hcro4- heptanoic acid(aq)	hbro(aq) hclo(aq) hcoo2- heptanoate hexanoate hf(aq) hio3(aq)
435 437 439 441 443 445 447 449	hass2(aq) hcl(aq) hclo2(aq) hcro4- heptanoic acid(aq) hexanoic acid(aq) hf2- hn3(aq)	hbro(aq) hclo(aq) hcoo2- heptanoate hexanoate hf(aq)
435 437 439 441 443 445 447 449 451	hass2(aq) hcl(aq) hclo2(aq) hcro4- heptanoic acid(aq) hexanoic acid(aq) hf2-	hbro(aq) hclo(aq) hcoo2- heptanoate hexanoate hf(aq) hio3(aq) hno2(aq)
435 437 439 441 443 445 447 449 451 453	hass2(aq) hcl(aq) hclo2(aq) hcro4- heptanoic acid(aq) hf2- hn3(aq) hno3(aq)	hbro(aq) hclo(aq) hcoo2- heptanoate hexanoate hf(aq) hio3(aq) hno2(aq) ho2-
435 437 439 441 443 445 447 449 451 453 455 457 459	hass2(aq) hcl(aq) hclo2(aq) hcro4- heptanoic acid(aq) hf2- hn3(aq) hno3(aq) hp2o7 hruo5- hse-	hbro(aq) hclo(aq) hcoo2- heptanoate hexanoate hf(aq) hio3(aq) hno2(aq) ho2- hpo3f- hs203- hseo3-
435 437 439 441 443 445 447 449 451 453 455 457 459 461	hass2(aq) hcl(aq) hclo2(aq) hcro4- heptanoic acid(aq) hf2- hn3(aq) hn03(aq) hp207 hru05- hse- hseo4-	hbro(aq) hclo(aq) hcoo2- heptanoate hexanoate hf(aq) hio3(aq) hno2(aq) hno2(aq) hoo2- hpo3f- hs2o3- hseo3- hsio3-
$\begin{array}{r} 435\\ 437\\ 439\\ 441\\ 443\\ 445\\ 445\\ 445\\ 453\\ 455\\ 457\\ 459\\ 461\\ 463\\ \end{array}$	hass2(aq) hcl(aq) hclo2(aq) hcro4- heptanoic acid(aq) hf2- hn3(aq) hno3(aq) hp2o7 hruo5- hse- hseo4- hso3-	hbro(aq) hclo(aq) hcoo2- heptanoate hexanoate hf(aq) hio3(aq) hno2(aq) hc2- hpo3f- hs2o3- hseo3- hsio3- hso4-
$\begin{array}{r} 435\\ 437\\ 439\\ 441\\ 443\\ 445\\ 445\\ 445\\ 453\\ 455\\ 457\\ 459\\ 461\\ 463\\ 465\\ \end{array}$	hass2(aq) hcl(aq) hclo2(aq) hcro4- heptanoic acid(aq) hf2- hn3(aq) hn03(aq) hp207 hru05- hse- hseo4- hso3- hso5-	hbro(aq) hclo(aq) hcoo2- heptanoate hexanoate hf(aq) hio3(aq) hno2(aq) ho2- hpo3f- hs2o3- hseo3- hsio3- hso4- hvo4
$\begin{array}{r} 435\\ 437\\ 439\\ 441\\ 443\\ 445\\ 447\\ 453\\ 455\\ 455\\ 457\\ 463\\ 465\\ 467\\ \end{array}$	hass2(aq) hcl(aq) hclo2(aq) hcro4- heptanoic acid(aq) hf2- hn3(aq) hn03(aq) hp207 hru05- hse- hseo4- hso3- hso5- i3-	hbro(aq) hclo(aq) hcoo2- heptanoate hexanoate hf(aq) hio3(aq) hno2(aq) hno2(aq) hno3f- hs203- hseo3- hsio3- hso4- hvo4 io-
$\begin{array}{r} 435\\ 437\\ 439\\ 441\\ 445\\ 445\\ 445\\ 455\\ 455\\ 455\\ 466\\ 365\\ 466\\ 466\\ 9\end{array}$	hass2(aq) hcl(aq) hclo2(aq) hcro4- heptanoic acid(aq) hf2- hn3(aq) hno3(aq) hp2o7 hruo5- hse- hseo4- hso3- hso5- i3- io4-	hbro(aq) hclo(aq) hcoo2- heptanoate hexanoate hf(aq) hio3(aq) hno2(aq) ho2- hpo3f- hs2o3- hseo3- hso3- hso4- hvo4 io- isoleucine(aq)
$\begin{array}{r} 435\\ 437\\ 439\\ 441\\ 443\\ 445\\ 445\\ 451\\ 455\\ 455\\ 463\\ 465\\ 466\\ 469\\ 471 \end{array}$	hass2(aq) hcl(aq) hclo2(aq) hcro4- heptanoic acid(aq) hf2- hn3(aq) hno3(aq) hp2o7 hruo5- hse- hseo4- hso5- i3- io4- kbr(aq)	hbro(aq) hclo(aq) hcoo2- heptanoate hexanoate hf(aq) hno2(aq) hno2(aq) hc2- hpo3f- hseo3- hsio3- hso4- hvo4 io- isoleucine(aq) kcl(aq)
$\begin{array}{r} 435\\ 437\\ 443\\ 444\\ 445\\ 445\\ 455\\ 455\\ 463\\ 465\\ 466\\ 466\\ 466\\ 473\\ \end{array}$	hass2(aq) hcl(aq) hclo2(aq) hcro4- heptanoic acid(aq) hf2- hn3(aq) hn207 hru05- hse- hse04- hso3- hso5- i3- io4- kbr(aq) khp04-	hbro(aq) hclo(aq) hcoo2- heptanoate hexanoate hf(aq) hio3(aq) hno2(aq) hno2(aq) hno2(aq) hso3- hso3- hsio3- hsio3- hso4- hvo4 isoleucine(aq) kcl(aq) khso4(aq)
$\begin{array}{r} 435\\ 437\\ 443\\ 443\\ 444\\ 445\\ 445\\ 455\\ 455\\ 465\\ 465\\ 465$	hass2(aq) hcl(aq) hclo2(aq) hcro4- heptanoic acid(aq) hf2- hn3(aq) hno3(aq) hp2o7 hruo5- hse- hso3- hso5- i3- io4- kbr(aq) khpo4- ki(aq)	hbro(aq) hclo(aq) hcoo2- heptanoate hexanoate hf(aq) hno2(aq) ho2- hpo3f- hs203- hse03- hse03- hso4- hvo4 io- isoleucine(aq) kcl(aq) kch(aq)
$\begin{array}{r} 435\\ 437\\ 441\\ 444\\ 445\\ 445\\ 455\\ 455\\ 465\\ 466\\ 467\\ 135\\ 467\\ 135\\ 7\\ 477\\ 7\end{array}$	hass2(aq) hcl(aq) hclo2(aq) hcro4- heptanoic acid(aq) hf2- hn3(aq) hno3(aq) hp2o7 hruo5- hse- hseo4- hso5- i3- io4- kbr(aq) khpo4- ki(aq) kp207	hbro(aq) hclo(aq) hcoo2- heptanoate hexanoate hf(aq) hno2(aq) hno2(aq) hc2- hso3- hsio3- hsio3- hso4- hvo4 io- isoleucine(aq) kcl(aq) kso4(aq) kso4-
$\begin{array}{c} 435\\ 437\\ 441\\ 444\\ 444\\ 445\\ 445\\ 445\\ 445\\ 445$	hass2(aq) hcl(aq) hclo2(aq) hcro4- heptanoic acid(aq) hf2- hn3(aq) hn207 hru05- hse- hse04- hso3- hso5- i3- i04- kbr(aq) khp04- ki(aq) kp207 leucine(aq)	hbro(aq) hclo(aq) hcoo2- heptanoate hexanoate hf(aq) hio3(aq) hno2(aq) hno2(aq) hno2(aq) hso3- hso3- hsio3- hso4- hvo4 isoleucine(aq) kcl(aq) khso4(aq) kso4- licl(aq)
$\begin{array}{c} 435\\ 437\\ 441\\ 444\\ 445\\ 445\\ 455\\ 455\\ 455\\ 461\\ 465\\ 465\\ 477\\ 477\\ 477\\ 481\\ \end{array}$	hass2(aq) hcl(aq) hclo2(aq) hcro4- heptanoic acid(aq) hf2- hn3(aq) hno3(aq) hp2o7 hruo5- hse- hso3- hso5- i3- i04- kbr(aq) khp04- ki(aq) kp2o7 leucine(aq) lioh(aq)	hbro(aq) hclo(aq) hcoo2- heptanoate hexanoate hf(aq) hno2(aq) ho2- hpo3f- hs203- hseo3- hseo3- hso4- hvo4 io- isoleucine(aq) kcl(aq) kch(aq) kso4- licl(aq) liso4-
$\begin{array}{c} 435\\ 4437\\ 444\\ 4445\\ 4451\\ 3557\\ 91\\ 3557\\ 91\\ 3557\\ 91\\ 3557\\ 91\\ 3557\\ 91\\ 3577\\ 91\\ 3577\\ 91\\ 33\\ 3577\\ 91\\ 33\\ 3577\\ 91\\ 33\\ 3577\\ 91\\ 33\\ 3577\\ 91\\ 33\\ 3577\\ 91\\ 33\\ 3577\\ 91\\ 3577$ 91\\ 3577 9	hass2(aq) hcl(aq) hclo2(aq) hcro4- heptanoic acid(aq) hf2- hn3(aq) hno3(aq) hp2o7 hruo5- hse- hseo4- hso3- hso5- i3- io4- kbr(aq) khpo4- ki(aq) kp2o7 leucine(aq) lioh(aq)	hbro(aq) hclo(aq) hcoo2- heptanoate hexanoate hf(aq) hno2(aq) hno2(aq) ho2- hpo3f- hseo3- hseo3- hseo3- hso4- hvo4 io- isoleucine(aq) kcl(aq) kcl(aq) kcb(aq) kco4- lic1(aq) liso4- mg(h3sio4)2(aq)
43579135791444455791355791355791355791355791355791355791355791355791355791355791355791355791355579135557913555791355579135557913555791355579135557991355557991355557991355557991355557991355579913555579913555799135555799135555799135555799135555799135555799135555799135555799135555799135555799135555555555	hass2(aq) hcl(aq) hclo2(aq) hcro4- heptanoic acid(aq) hf2- hn3(aq) hp207 hru05- hse- hse04- hso3- hso5- i3- io4- kbr(aq) khp04- ki(aq) kp207 leucine(aq) lioh(aq) mg4(oh)4++++	hbro(aq) hclo(aq) hcoo2- heptanoate hexanoate hf(aq) hio3(aq) hno2(aq) hno2(aq) hso3- hso3- hsio3- hso4- hvo4 io- isoleucine(aq) kcl(aq) khso4(aq) kch(aq) kso4- licl(aq) liso4- mg(h3sio4)2(aq) mgb(oh)4+
$\begin{array}{c} 435\\ 4437\\ 444\\ 4445\\ 4451\\ 3557\\ 91\\ 3557\\ 91\\ 3557\\ 91\\ 3557\\ 91\\ 3557\\ 91\\ 3577\\ 91\\ 3577\\ 91\\ 33\\ 3577\\ 91\\ 33\\ 3577\\ 91\\ 33\\ 3577\\ 91\\ 33\\ 3577\\ 91\\ 33\\ 3577\\ 91\\ 33\\ 3577\\ 91\\ 3577$ 91\\ 3577 9	hass2(aq) hcl(aq) hclo2(aq) hcro4- heptanoic acid(aq) hf2- hn3(aq) hno3(aq) hp207 hru05- hse- hse04- hso3- hso5- i3- i04- kbr(aq) khp04- ki(aq) kp207 leucine(aq) lioh(aq) mg4(oh)4++++ mgch3coo+	hbro(aq) hclo(aq) hcoo2- heptanoate hexanoate hf(aq) hno2(aq) hno2(aq) ho2- hpo3f- hseo3- hseo3- hseo3- hso4- hvo4 io- isoleucine(aq) kcl(aq) kcl(aq) kcb(aq) kco4- lic1(aq) liso4- mg(h3sio4)2(aq)
435 437 439 441 443 445 447 449 451 455 457 461 463 465 467 469 471 467 467 477 477 481 485 485	hass2(aq) hcl(aq) hclo2(aq) hcro4- heptanoic acid(aq) hf2- hn3(aq) hno3(aq) hp2o7 hruo5- hse- hseo4- hso3- hso5- i3- i04- kbr(aq) khp04- ki(aq) kp2o7 leucine(aq) lioh(aq) methionine(aq) mg4(oh)4++++ mgch3coo+ mgco3(aq)	hbro(aq) hclo(aq) hcoo2- heptanoate hexanoate hf(aq) hno2(aq) hno2- hpo3f- hso3- hso3- hso3- hso4- hvo4 isoleucine(aq) kcl(aq) kch(aq) kco4(aq) kco4- lic1(aq) liso4- mg(h3sio4)2(aq) mgb(ch)4+ mgc1+ mgf+
435 437 449 441 443 445 445 445 445 445 455 455 455 463 465 467 471 473 477 479 483 485 485	hass2(aq) hcl(aq) hclo2(aq) hcro4- heptanoic acid(aq) hf2- hn3(aq) hno3(aq) hp207 hru05- hse- hse04- hso3- hso5- i3- i04- kbr(aq) khp04- ki(aq) kp207 leucine(aq) lioh(aq) mg4(oh)4++++ mgch3coo+	hbro(aq) hclo(aq) hcoo2- heptanoate hexanoate hf(aq) hno2(aq) hno2(aq) ho2- hgo3f- hseo3- hseo3- hseo3- hso4- hvo4 isoleucine(aq) kcl(aq) kcb(aq) kso4(aq) kso4(aq) kso4(aq) kso4- lic1(aq) liso4- mg(h3sio4)2(aq) mgb(ch)4+ mgcl+
435 437 449 443 445 445 445 445 445 445 445 445 445	<pre>hass2(aq) hcl(aq) hclo2(aq) hcro4- heptanoic acid(aq) hf2- hn3(aq) hp207 hru05- hse- hse04- hso3- hso5- i3- io4- kbr(aq) khp04- ki(aq) khp04- ki(aq) methionine(aq) methionine(aq) methionine(aq) mg4(oh)4++++ mgch3coo+ mgch3(aq) mgh2p04+</pre>	hbro(aq) hclo(aq) hcoo2- heptanoate hexanoate hf(aq) hio3(aq) hno2(aq) hno2(aq) hno2(aq) hso3- hso3- hso3- hso4- hvo4 io- isoleucine(aq) kcl(aq) khso4(aq) kch(aq) kso4- licl(aq) liso4- mg(h3sio4)2(aq) mgb(oh)4+ mgc1+ mgf+ mgf2sio4(aq)
435 437 449 443 445 445 445 445 445 445 445 445 445	hass2(aq) hcl(aq) hclo2(aq) hcro4- heptanoic acid(aq) hf2- hn3(aq) hno3(aq) hp207 hru05- hse- hse04- hso3- hso5- i3- i04- kbr(aq) khp04- ki(aq) kp207 leucine(aq) lioh(aq) mg4(oh)4++++ mgch3co0+ mgh2p04+ mgh3sio4+	hbro(aq) hclo(aq) hcco2- heptanoate hexanoate hf(aq) hio3(aq) hno2(aq) hno2(aq) hno2(aq) hso3- hso3- hsio3- hso4- hvo4 isoleucine(aq) kcl(aq) kcl(aq) kcd(aq) kcso4- lic1(aq) liso4- mg(h3sio4)2(aq) mgb(ch)4+ mgf+ mgf+ mgh2sio4(aq) mghco3+ mgh04(aq) mgbo04-
435 437 449 443 445 445 445 447 447 445 455 465 465 465 465 465 465 467 471 473 475 481 485 487 485 487 485 487 485 487 485 487 485 487 485 487 485 487 499 485 499 499 493	hass2(aq) hcl(aq) hclo2(aq) hcro4- heptanoic acid(aq) hf2- hn3(aq) hno3(aq) hp207 hruo5- hse- hse04- hso3- hso5- i3- i04- kbr(aq) khp04- ki(aq) kp207 leucine(aq) lioh(aq) methionine(aq) mg4(oh)4++++ mgch3co0+ mgco3(aq) mgh2p04- mgh2p07 mgp207 mgp207 mgp207 mgp207 mgp207 mgp207 mgp207 mgp207 mgp207 mgs04(aq)	hbro(aq) hclo(aq) hcco2- heptanoate hexanoate hf(aq) hio3(aq) hno2(aq) ho2- hgo3f- hsco3- hsco3- hsco3- hso4- hvo4 io- isoleucine(aq) kcl(aq) kcl(aq) kso4(aq) kso4(aq) kso4(aq) kso4(aq) kso4(aq) mgb(oh)4+ mgcl+ mgft2 mghco3+ mghco3+ mgpo4- m(no3)2(aq)
435 437 449 443 445 445 445 445 445 455 455 455 467 455 467 471 473 463 465 467 471 473 477 479 481 483 485 487 493 495 495 495 495 501	hass2(aq) hcl(aq) hclo2(aq) hcro4- heptanoic acid(aq) hf2- hn3(aq) hno3(aq) hp2o7 hruo5- hse- hso3- hso5- i3- i04- kbr(aq) khp04- ki(aq) kp2o7 leucine(aq) lioh(aq) methionine(aq) mg4(oh)4++++ mgch3coo+ mgco3(aq) mgh2p04+ mgh2p07 mgp2o7 mgp2o7 mgp2o7 mgp2o7 mgs04(aq) mn(oh)2(aq)	hbro(aq) hclo(aq) hclo(aq) hcoo2- heptanoate hexanoate hf(aq) hno2(aq) ho2- hpo3f- hs203- hse03- hse03- hso4- hvo4 isoleucine(aq) kcl(aq) kch(aq) kso4(aq) kco4(aq) kso4- licl(aq) liso4- mg(h3sio4)2(aq) mgb(ch)4+ mgf+ mgf2sio4(aq) mgbc3+ mghpo4(aq) mgpo4- mn(no3)2(aq) mn(ch)3-
435 437 449 443 445 445 445 445 445 445 445 445 445	hass2(aq) hcl(aq) hclo2(aq) hcro4- heptanoic acid(aq) hf2- hn3(aq) hp207 hru05- hse- hse04- hso3- hso5- i3- i04- kbr(aq) khp04- ki(aq) khp04- ki(aq) methionine(aq) methionine(aq) methionine(aq) mg4(oh)4++++ mgch3co0+ mg207 mgp207	hbro(aq) hclo(aq) hcco2- heptanoate hexanoate hf(aq) hio3(aq) hno2(aq) hno2(aq) hoo2- hpo3f- hsco3- hsio3- hso4- hvo4 io- isoleucine(aq) kcl(aq) khso4(aq) khso4(aq) kso4- licl(aq) liso4- mg(h3sio4)2(aq) mgb(ch)4+ mgc1+ mgf+ mghc3+ mghpo4(aq) mgbc03- mgbo4- mn(no3)2(aq) mn(ch)3- mn2(ch)3+
435 437 439 441 443 445 445 447 449 451 455 465 465 465 465 465 465 465 467 471 473 475 481 485 485 485 485 485 491 493 495 503 503 503	hass2(aq) hcl(aq) hclo2(aq) hcro4- heptanoic acid(aq) hf2- hn3(aq) hp207 hru05- hse- hse04- hso3- hso5- i3- i04- kbr(aq) khp04- ki(aq) kp207 leucine(aq) lioh(aq) methionine(aq) mg4(oh)4+++ mgch3co4- mg207 mgp207	hbro(aq) hclo(aq) hcoo2- heptanoate hexanoate hf(aq) hio3(aq) hno2(aq) hoo2- hpo3f- hseo3- hseo3- hso3- hso4- hvo4 isoleucine(aq) kcl(aq) kcl(aq) kso4(aq) kbs04(aq) kso4- lic1(aq) liso4- mg(h3sio4)2(aq) mgb(oh)4+ mgft+ mgft+ mgfx(aq) mgbo4(aq) mgbo4(aq) mgbo4(aq) mgbo4- mn(no3)2(aq) mn(oh)3- mn2(oh)3+ mncl+
435 437 449 443 445 445 445 445 445 455 457 459 461 455 467 471 473 463 465 467 471 473 477 479 481 483 487 489 491 495 495 497 9501 505 505 507	hass2(aq) hcl(aq) hclo2(aq) hcro4- heptanoic acid(aq) hf2- hn3(aq) hno3(aq) hp2o7 hru5- hse- hse- hso5- i3- i04- kbr(aq) khp04- ki(aq) kp2o7 leucine(aq) lioh(aq) mg4(oh)4++++ mgch3coo+ mgco3(aq) mgh2o7 mgp2o7	hbro(aq) hclo(aq) hcoo2- heptanoate hexanoate hf(aq) hno2(aq) hno2- hpo3f- hs203- hseo3- hseo3- hseo3- hso4- hvo4 io- isoleucine(aq) kcl(aq) kch(aq) kch(aq) kso4- licl(aq) liso4- mg(h3sio4)2(aq) mgb(ch)4+ mgf+ mgf2+ mgf2sio4(aq) mgbo4- mgh03- hso3- hso4- ngf(ag) hso4- hog12(aq) mgp04- mn(no3)2(aq) mnch)3- mn2(ch)3+ mnch+ mnco3(aq)
435 437 443 443 444 443 445 445 445 445 445 445	hass2(aq) hcl(aq) hclo2(aq) hcro4- heptanoic acid(aq) hf2- hn3(aq) hp207 hru05- hse- hse04- hso3- hso5- i3- i04- kbr(aq) khp04- ki(aq) khp04- ki(aq) kp207 leucine(aq) lioh(aq) mg4(oh)4++++ mgch3co0+ mg4207 mgp20	hbro(aq) hclo(aq) hcoo2- heptanoate hexanoate hf(aq) hio3(aq) hno2(aq) hno2(aq) hno2(aq) hso3- hso3- hso3- hso4- hso4- hso4- isoleucine(aq) kcl(aq) khso4(aq) khso4(aq) kso4- lic1(aq) liso4- mg(h3sio4)2(aq) mgb(ch)4+ mgc1+ mgf+ mghc3+ mghpo4(aq) mghco3+ mghpo4(aq) mn(oh)3- mn(oh)3- mn(oh)3+ mnc03(aq) mh2po4+
435 437 441 443 445 445 445 445 445 445 445 445 445	hass2(aq) hcl(aq) hclo2(aq) hcro4- heptanoic acid(aq) hf2- hn3(aq) hn03(aq) hp207 hru05- hse- hse04- hso3- hso5- i3- i04- kbr(aq) khp04- ki(aq) kp207 leucine(aq) lioh(aq) methionine(aq) mg4(oh)4+++ mgch3co4- mg520(aq) hoftaq mg4207 mg520	hbro(aq) hclo(aq) hcoo2- heptanoate hexanoate hf(aq) hio3(aq) hno2(aq) hno2(aq) hoo2- hpo3f- hseo3- hseo3- hso3- hso4- hvo4 isoleucine(aq) kcl(aq) kcl(aq) kso4(aq) kso4(aq) kso4- licl(aq) liso4- mg(h3sio4)2(aq) mgb(oh)4+ mgft+ mgf2+ mgho4(aq) mgbo4(aq) mn(oh)3- mn2(oh)3+ mncl+ mncl2(aq) mh2po4+ mnh2po4+ mnh2po4+ mnh2po4+ mnh2po4(aq)
435 437 441 443 445 445 445 445 445 445 465 465 465 465	hass2(aq) hcl(aq) hclo2(aq) hcro4- heptanoic acid(aq) hf2- hn3(aq) hp207 hru05- hse- hse04- hso3- hso5- i3- i04- kbr(aq) khp04- ki(aq) khp04- ki(aq) kp207 leucine(aq) lioh(aq) mg4(oh)4++++ mgch3co0+ mg4207 mgp20	hbro(aq) hclo(aq) hcoo2- heptanoate hexanoate hf(aq) hio3(aq) hno2(aq) hno2(aq) hno2(aq) hso3- hso3- hso3- hso4- hso4- hso4- isoleucine(aq) kcl(aq) khso4(aq) khso4(aq) kso4- lic1(aq) liso4- mg(h3sio4)2(aq) mgb(ch)4+ mgc1+ mgf+ mghc3+ mghpo4(aq) mghco3+ mghpo4(aq) mn(oh)3- mn(oh)3- mn(oh)3+ mnc03(aq) mh2po4+

517	mnseo4(aq)
519	n-butane(aq)
521	n-heptane(aq)
523	n-hexane(aq)
525	n-octane(aq)
527 529	n-pentane(aq)
531	n-propylbenzene(aq) na2p2o7
533	nabr(aq)
535	nacl(aq)
537	naf(aq)
539	nahco3(aq)
541	nahpo4-
543	nai(aq)
545	nap2o7
547	nh3(aq)
549	ni(nh3)2++
551	ni(no3)2(aq)
553	ni(oh)3-
555	ni4(oh)4++++
557	nicl+
559	nino3+
561	niseo4(aq)
563	np(co3)5(6-)
565	np(h2po4)3(aq)
567	np(hpo4)3
569	np(hpo4)5(6-)
571	np(oh)3+
573	np(oh)5-
575	npcl+++
577	npf+++
579	nph2po4++
581	npo2(co3)2
583 585	npo2(co3)3(5-)
587	npo2cl(aq) npo2co3-
589	npo2f+
591	npo2h2po4(aq)
593	npo2hpo4(aq)
595	npo2oh(aq)
597	npo2so4(aq)
599	npoh++
601 603	npso4++
605	octanoic acid(aq) pb(ch3coo)2(aq)
607	pbcl+
609	pbcl3-
611	pbf+
613	pbh2po4+
615	pentanoate
617	ph4+
619	phenylalanine(aq)
621	propane(aq)
623	propanoic acid(aq)
625	pu(hpo4)3
627	pu(oh)2++
629	pu(oh)4(aq)
631	pu(so4)2(aq)
633	pucl++
635	puf+++
637	puf3+
639	puh2po4++
641	puo2(co3)2
643	puo2f+
645	puo2f3-
647	puo2h2po4+
649	puo2oh+
651	puoh++
653	puso4+
655	rbbr(aq)
657	rbf(aq)
659	ru(cl)2+
661	ru(oh)2+
663	ru(oh)2cl2(aq)
665	ru(oh)2cl4
667	ru(oh)4(aq)
669	ru4(oh)12++++
671	rucl++
673	rucl5
675	ruoh++
677	ruso4+
679	s2
681	s2o4
683	s2o6
685	s3
687	s4
689	s5
691	scn-
693	sif6
695	sn(oh)2++
697	sn(oh)3-
699	sn(so4)2(aq)
701	sncl2(aq)

mnso4(aq)
n-butylbenzene(aq)
n-heptylbenzene(aq)
n-hexylbenzene(aq)
n-octylbenzene(aq) n-pentylbenzene(aq) na(o-phth)-nab(oh)4(aq) nach3coo(aq) naco3nah3sio4(aq) nahp2o7--nahsio3(aq) naoh(aq) naso4-nh4so4ni(nh3)6++ ni(oh)2(aq) ni2oh+++ ni2on+++ nibr+ nihp2o7--nip2o7--niso4(aq) np(hpo4)2+ np(hpo4)2(aq) np(hpo4)4---np(oh)2++np(oh)4(aq)np(so4)2(aq)npcl2++ npf2++ nphpo4++ npo2(co3)2---npo2(co3)3---npo2cl+ npo2f(aq) npo2f2(aq) npo2h2po4+ npo2hpo4npo2oh+ npo2so4npoh+++ octanoate p207----pbch3coo+ pbcl2(aq) pbcl4--pbf2(aq) pbhpo4(aq) pbnpo4(aq)
pentancic acid(aq)
phenol(aq)
po3f-propanoate
pu(hpo4)2(aq)
pu(hpo4)4---pu(oh)3+ pu(oh)3+ pu(oh)5-pu(so4)2pucl+++ puf2++ puf4(aq) puhpo4++ puo2cl+ puo2f2(aq) puo2f4-puo2oh(aq) puo2so4(aq) puoh+++ puso4++ rbcl(aq) rbi(aq) ru(cl)3(aq) ru(oh)2cl+ ru(oh)2cl3ru(oh)2so4(aq) ru(so4)2rucl+ rucl4rucl6---ruso4(aq) s--s2o3-s2o5-s208-s306--s406-s506-serine(aq)
sn(oh)2(aq)
sn(oh)3+ sn(oh)4(aq) sncl+ sncl3-

703 705 707	snf+ snf3- snoh+++	snf2(aq) snoh+ snso4++
	so2(aq)	srch3coo+
711	srcl+	srco3(aq)
	srf+	srh2po4+
715 717	srhpo4(aq) sroh+	srno3+ srp2o7
719	srpo4-	srso4(aq)
721	tc++	tco(oh)2(aq)
723	tcooh+	th(h2po4)2++
725 727	th(hpo4)2(aq) th(oh)2++	th(hpo4)3 th(oh)4(aq)
729	th(so4)2(aq)	th(so4)3
731	th(so4)4	th2(oh)2(6+)
733 735	th4(oh)8(8+) thcl+++	th6(oh)15(9+) thcl2++
737	thcl3+	thcl4(aq)
739	thf+++	thf2++
741 743	thf3+	thf4(aq)
745	thh2po4+++ thoh+++	thh3po4++++ threonine(aq)
747	thso4++	tl+++
749	toluene(aq)	tryptophan(aq)
751 753	tyrosine(aq) u(co3)5(6-)	u(co3)4 u(no3)2++
755	u(oh)4(aq)	u(scn)2++
757	u(so4)2(aq)	ubr+++
759	ucl+++	uf+++
761 763	uf2++ uf4(aq)	uf3+ uf5-
765	uf6	ui+++
767	uno3+++	uo2(co3)2
769 771	uo2(co3)3(5-) uo2(h2po4)2(aq)	uo2(co3)3 uo2(h2po4)h3po4+
773	uo2(io3)2(aq)	uo2(n3)2(aq)
775	uo2(n3)3-	uo2(n3)4
777 779	uo2(oh)2(aq) uo2(oh)4	uo2(oh)3-
781	uo2(scn)3-	uo2(scn)2(aq) uo2(so3)2
783	uo2(so4)2	uo2br+
785 787	uo2bro3+	uo2cl+
789	uo2cl2(aq) uo2co3(aq)	uo2clo3+ uo2f+
791	uo2f2(aq)	uo2f3-
793	uo2f4	uo2h2po4+
795 797	uo2h3po4++ uo2io3+	uo2hpo4(aq) uo2n3+
799	uo2no3+	uo2oh+
801	uo2po4-	uo2s2o3(aq)
803 805	uo2scn+ uo2so4(aq)	uo2so3(aq) uoh+++
807	uscn+++	uso4++
809	v(oh)2+	v2(oh)2++++
811 813	valine(aq) vo2(hpo4)2	vo(oh)3(aq) vo2(oh)2-
815	vo2f(aq)	vo2f2-
817	vo2h2po4(aq)	vo2hpo4-
819 821	vo2so4- vof+	vo3oh vof2(aq)
823	vol++	vooh+
825	voso4(aq)	vso4+
827	zn(ch3coo)2(aq)	zn(ch3coo)3-
	znch3coo+ zncl2(ag)	zncl+ zncl3-
833	zncl2(aq) zncl4	znf+
	znh2po4+	znhco3+
	znhpo4(aq) znpo4-	znoh+ znseo4(aq)
	zr(oh)3+	zr(oh)4(aq)
	zr(so4)2(aq)	zr(so4)3
845 847	zrf+++ zrf3+	zrf2++ zrf4(aq)
849		zrf6
	zroh+++	zrso4++
minera	ls	
1	(pb(oh)2)3.pbcl2	(uo2)2as2o7
3	(uo2)2cl3	(uo2)2aS207 (uo2)2p2o7
5	(uo2)3(aso4)2	(uo2)3(po4)2
7 9	(uo2)3(po4)2:4h2o acanthite	(vo)3(po4)2 afwillite
9 11	silver	ag3po4
13	ahlfeldite	akermanite
15	al	al2(so4)3
17 19	al2(so4)3:6h2o alamosite	alabandite albite
21	albite high	albite low
23 25	alf3 alum-k	alstonite alunite
25	am	am(oh)3

29	am(oh)3(am)
31	amohco3
33	analcime-dehy
35	andradite
37	anhydrite
39	anorthite
41 43	anthophyllite
45	antlerite aragonite
47	arsenolite
49	artinite
51	as2o5
53	as4o6(octa)
55	gold
57	b
59	ba
61	ba2si3o8
63	babr2
65	bacl2
67	bacl2:h2o
69	baddeleyite
71	bai2
73	bao
75	barytocalcite
77	baseo3
79	basif6
81	bassetite
83	bazro3
85	bel3u
87	beidellite-cs
89	beidellite-k
91 93	beidellite-na
95	berndtite birnessite
97	bixbyite
99	boehmite
101	boltwoodite-na
103	boric acid
105	brezinaite
107	brucite
109	bunsenite
111	graphite
113	ca-al pyroxene
115 117	ca2cl2(oh)2:h2o
119	ca3(aso4)2 ca3v2o8
121	ca4al2o7:13h2o
123	ca4cl2(oh)6:13h2o
125	caal2o4:10h2o
127	cadmoselite
129	calomel
131	carnotite
133	caseo4
135	cassiterite
137	cauo4
139	cazro3
141	cdcr2o4
143	cdseo4
145	ce
147	celestite
149	chalcanthite
151	chalcocite
153	chalcopyrite
155	chlorargyrite
157	chromite
159	chrysotile
161	claudetite
163	clinochalcomenite
165	clinochlore-7a
167	clinoptilolite-ca
169	clinoptilolite-dehy
171	clinoptilolite-dehy-cs
173	clinoptilolite-dehy-na
175	clinoptilolite-dehy-sr
177	clinoptilolite-na
179	clinoptilolite-sr
181	co
183	co(oh)2
185	co3(aso4)2
187	cocl2
189	cocl2:6h2o
191	cof2
193	cofe2o4
195	cohpo4
197	coo
199	cordierite hydr
201	corundum
203	coseo3
205	coso4.3co(oh)2
207	coso4:h2o
209	covellite
211	cr
213	crf3
	-

amesite-14a analcime andalusite anglesite annite antarcticite antigorite aphthitalite arcanite arsenopyrite as as4o6(mono) atacamite azurite b2o3 ba(oh)2:8h2o ba2sio4 babr2:2h2o bacl2:2h2o bacro4 bahpo4 bamno4 barite bas baseo4 bassanite bauo4 be beidellite-ca beidellite-h beidellite-mg berlinite bieberite bischofite bloedite boltwoodite borax bornite brochantite brushite burkeite ca ca2al2o5:8h2o ca2v2o7 ca3a12o6 ca4al2fe2o10 ca4al2o7:19h2o caal2o4 caal4o7 calcite carnallite caseo3:2h2o caso4:0.5h2o(beta) cattierite cav2o6 cd cd cdseo3 cdso4:8/3h2o celadonite cerussite chalcedony chalcocyanite chamosite-7a chloromagnesite chrysocolla cinnabar clausthalite clinochlore-14a clinoptilolite clinoptilolite-cs clinoptilolite-dehy-ca clinoptilolite-dehy-k clinoptilolite-dehy-h4 clinoptilolite-k clinoptilolite-h4 clinoptilolite-nh4 clinozisite co(no3)2 co2sio4 co3(po4)2 cocl2:2h2o coesite cof3 coffinite colemanite cordierite anhyd corkite cos coso4 coso4:6h2o cotunnite cowo4 crcl3 crf4

215	cri3	C
217	cristobalite-a	C
219	cro2	C
221	crocoite	C
223	crs	C
225	cs2u2o7	C
227	copper	C.
229	cu3(po4)2:3h2o	C.
231	cucr2o4	C.
233	cuf2	C.
235	cuprite	C.
237	daphnite-14a	d
239	dawsonite	d
241	diaspore	d
243	diopside	d
245	dolomite	d
247	dolomite-ord	d
249 251	dy amidata	e
253	epidote epsomite	ej
255	erythrite	e
257	ettringite	e
259	eu(io3)3:2h2o	e
261	eu(oh)2.5cl.5	e
263	eu(oh)3	e
265	eu2(so4)3:8h2o	e
267	eu2o3(monoclinic)	e
269	eubr3	e
271	eucl3	e
273	eucryptite	e
275	euo	e
277	eus	e
279	fayalite	i
281	fe(oh)2	f
283	fe2(so4)3 fef2	f
285 287	feo	f
289	ferrite-cu	f
291	ferrite-mg	f
293	ferroselite	f
295	feso4	f
297	fluorapatite	f
299	forsterite	f
301	frankdicksonite	f
303	ga	g
305	gaylussite	g
307	gehlenite	g
309	gismondine	g
311	goethite	g
313 315	grossular	g h
317	gyrolite haiweeite	h
319	hatrurite	h
321	heazlewoodite	h
323	hematite	h
325	herzenbergite	h
327	hexahydrite	h
329	hg2so4	h
331	hillebrandite	h
333	ho	h
335	htco4	h
337	hydroboracite	h
339	hydromagnesite	h
341	hydroxylapatite	i
343	ice	i
345 347	ilmenite jadeite	i: j
349	jarosite-na	J. k
351	k-feldspar	k
353	k2o	k
355	k2uo4	k
357	k8h4(co3)6:3h2o	k
359	kal(so4)2	k
361	kalsilite	k
363	karelianite	k
365	katoite	k
367	kieserite	k
369	kmgcl3	ki
371	knaco3:6h2o	k
373	ktco4	k
375 377	kyanite lammerite	1.
379	lansfordite	1
381	laumontite	1
383	lawrencite	1.
385	leonite	1
387	li2se	1
389	lime	ī
391	litharge	1
393	lopezite	1
395	magnesiochromite	m
397	magnetite	m
399	manganite	m

cristobalite cristobalite-b ro3 cronstedtite-7a s cs2u4o12 cu3(po4)2 cuc12 uf 2uf2:2h2o cuf2:2n2o cuseo3 laphnite-7a lelafossite dicalcium silicate lioptase lolomite-dis lowneyite enstatite epidote-ord er eskolaite eu eu eu(no3)3:6h2o eu(oh)2cl eu2(co3)3:3h2o eu2o3(cubic) eu3o4 eucl2 eucl3:6h2o euf3:0.5h2o euocl uso4 iron Ee(oh)3 easo4 fef3 ferrite-ca ferrite-dicalcium ferrite-zn errosilite ev2o4 luorite oshagite reboldite galena galena gd gibbsite glauberite greenalite yypsum 1-autunite 1alite nalite hausmannite nedenbergite nercynite neulandite ng2seo3 1928e03 198e03 1insdalite 1opeite 1untite hydrocerussite hydrophilite i2 illite n jarosite 2co3:3/2h2o 2se 3h(so4)2 ainite alicinite aolinite asolite br lockmannite mgcl3:2h2o rutaite uo2aso4 a anarkite larnite laurite awsonite i2uo4 innaeite iuo2aso4 u nagnesite nalachite anganosite

$\begin{array}{c} 4444444444444444444$	<pre>margarite matlockite mayenite mercallite mesolite mg mg1.5so4(oh) mg3(aso4)2 mgb72:6h2o mgoh21 mgseo3:6h2o mgu04 millerite minesotaite misenite mn(oh)2(am) mn3(aso4)2 mncl2:2h2o mncl2:2h2o mncl2:2h2o mncl2:2h2o mncl2:2h2o mncl2:2h2o mncl2:2h2o mncl2:2h2o mncl2:may mseo3 mso4 mo molysite montmor-ca montmor-ca montmor-k montmor-na mordenite misenite muscovite na2co3 na2cr2o7 na2o na2se2 na2u2o7 na3h(so4)2 na4uc2(co3)3 nabr nafeo2 napo2co3:3.5h2o natco4 natron naumannite nd nesquehonite ni ni2p2o7 ni3(po4)2 nicl2 nicl2:4h2o nif2 ningyoite nontronite-na np(hpo4)2 np2o5 npo2(oh)2 o-phthalic acid orpiment ottemannite p paralaurionite parsonsite pb(h2po4)2 pb40(po4)2 pb3(po4)2 pb40(po4)2 pb40(po4)2</pre>
583	pyrophyllite
585	quartz

massicot maximum microcline melanterite merwinite merwinite
metacinnabar
mg1.25so4(oh)0.5:0.5h2o
mg2v2o7
mg5r2
mgc12:2h2o
mgc12:h2o
mgseo3
mgca4 mgso4 mgv2o6 minium mirabilite mn mn(oh)3 mn3(po4)2 mncl2:4h2o mnhpo4 mnse mnseo3:2h2o mnv2o6 modderite monohydrocalcite monticellite montmor-cs montmor-mg montroydite mordenite-dehy mose2 na na2co3:7h2o na2cro4 na2se na2sio3 na2uo4(alpha) na3uo4 na4sio4 na6si2o7 nabr:2h2o nahcolite nantokite natrolite natrosilite nauo3 nepheline nh4hse ni(oh)2 ni2sio4 nickelbischofite nicl2:2h2o nico3 nif2:4h2o niso4 niter nontronite-ca nontronite-h nontronite-mg np np(oh)4 npo2 npo2oh(am) okenite otavite oxychloride-mg paragonite pargasite pb pb2sio4 pb3so6 pb4so7 pbf2 pbseo4 pd pentahydrite petalite phosgenite pirssonite plumbogummite polyhalite pr przhevalskite pu pu(oh)3 pu2o3(beta) puf4 puo2(oh)2 puo2oh(am) pyrolusite pyromorphite-oh pyrrhotite ra

587	ra(no3)2	racl2:2h2o
589	rankinite	raso4
591	rb	rb2uo4
593	re	realgar
595	rhodochrosite	rhodonite
597 599	ripidolite-14a romarchite	ripidolite-7a ru
601	ru(oh)3:h2o(am)	rubr3
603	rucl3	rui3
605	ruo2	ruo2:2h2o(am)
607	ruo4	rutherfordine
609	rutile	sulfur
611	safflorite	saleeite
613	sanbornite	sanidine high
615 617	saponite-ca saponite-h	saponite-cs saponite-k
619	saponite-mg	saponite-na
621	sc	scacchite
623	schoepite	<pre>schoepite-dehy(.393)</pre>
625	schoepite-dehy(.648)	<pre>schoepite-dehy(.85)</pre>
627	schoepite-dehy(.9)	schoepite-dehy(1.0)
629 631	scolecite se2o5	se secl4
633	sellaite	seo3
635	sepiolite	shcherbinaite
637	si	siderite
639	sillimanite	sio2(am)
	sklodowskite	sm
643	smectite-high-fe-mg	smectite-low-fe-mg
645	smithsonite	tin
647 649	sn(oh)2 sn3s4	sn(so4)2 snbr2
651	snbr4	sncl2
653	snse	snse2
655	snso4	soddyite
657	sphaerocobaltite	sphalerite
659	spinel	spinel-co
661	spodumene	sr (2) a. N. a
663	sr(no3)2	sr(no3)2:4h2o
665 667	sr(oh)2 sr3(aso4)2	sr2sio4 srbr2
669	srbr2:6h2o	srbr2:h2o
671	srcl2	srcl2:2h2o
673	srcl2:6h2o	srcl2:h2o
675	srcro4	srf2
677	srhpo4	sri2
679	sro	srs
681	srseo4	srsio3
683 685	sruo4(alpha)	srzro3
687	starkeyite stilleite	stilbite strengite
689	strontianite	sylvite
691	syngenite	tachyhydrite
693	talc	tarapacaite
695	tb	tc
697	tc(oh)2	tc(oh)3
699	tc2s7	tc304
701	tc407	tco2:2h2o(am)
703 705	tco3 tcs2	tcoh tcs3
707	tenorite	
709		
	th	tephroite th(no3)4:5h2o
711	th th(oh)4	
713	th(oh)4 th2s3	th(no3)4:5h2o th(so4)2 th2se3
713 715	th(oh)4 th2s3 th7s12	th(no3)4:5h2o th(so4)2 th2se3 thbr4
713 715 717	th(oh)4 th2s3 th7s12 thc14	th(no3)4:5h2o th(so4)2 th2se3 thbr4 thenardite
713 715 717 719	th(oh)4 th2s3 th7s12 thc14 thermonatrite	th(no3)4:5h2o th(so4)2 th2se3 thbr4 thenardite thf4
713 715 717 719 721	th(oh)4 th2s3 th7s12 thc14 thermonatrite thf4:2.5h2o	<pre>th(no3)4:5h2o th(so4)2 th2se3 thbr4 thenardite thf4 thi4</pre>
713 715 717 719	th(oh)4 th2s3 th7s12 thc14 thermonatrite	th(no3)4:5h2o th(so4)2 th2se3 thbr4 thenardite thf4 thi4 ths2
713 715 717 719 721 723	th(oh)4 th2s3 th7s12 thc14 thermonatrite thf4:2.5h2o thorianite	<pre>th(no3)4:5h2o th(so4)2 th2se3 thbr4 thenardite thf4 thi4</pre>
713 715 717 719 721 723 725	th(oh)4 th2s3 th7s12 thc14 thermonatrite thf4:2.5h2o thorianite ti	<pre>th(no3)4:5h2o th(so4)2 th2se3 thbr4 thenardite thf4 thi4 ths2 tiemannite</pre>
713 715 717 719 721 723 725 727 729 731	<pre>th(oh)4 th2s3 th7s12 thc14 thermonatrite thf4:2.5h2o thorianite ti titanite tm tobermorite-14a</pre>	<pre>th(no3)4:5h2o th(so4)2 th2se3 thbr4 thenardite thf4 thi4 ths2 tiemannite tl tobermorite-11a tobermorite-9a</pre>
713 715 717 719 721 723 725 727 729 731 733	<pre>th(oh)4 th2s3 th7s12 thc14 thermonatrite thf4:2.5h2o thorianite ti titanite tm tobermorite-14a todorokite</pre>	<pre>th(no3)4:5h2o th(so4)2 th2se3 thbr4 thenardite thf4 thi4 ths2 tiemannite tl tobermorite-11a tobermorite-9a torbernite</pre>
713 715 717 719 721 723 725 727 729 731 733 735	<pre>th(oh)4 th2s3 th7s12 thc14 thermonatrite thf4:2.5h2o thorianite ti titanite tm tobermorite-14a todorokite tremolite</pre>	<pre>th(no3)4:5h2o th(so4)2 th2se3 thbr4 thenardite thf4 thi4 ths2 tiemannite tl tobermorite-11a tobermorite-9a torbernite trevorite</pre>
713 715 717 719 721 723 725 727 729 731 733 735 737	<pre>th(oh)4 th2s3 th7s12 thc14 thermonatrite thf4:2.5h2o thorianite ti titanite tm tobermorite-14a todorokite tremolite tridymite</pre>	<pre>th(no3)4:5h2o th(so4)2 th2se3 thbr4 thenardite thf4 thi4 ths2 tiemannite tl tobermorite-11a tobermorite-9a torbernite trevorite troilite</pre>
713 715 717 719 721 723 725 727 729 731 733 735 737 739	<pre>th(oh)4 th2s3 th7s12 thc14 thermonatrite thf4:2.5h2o thorianite ti titanite tm tobermorite-14a todorokite tremolite tridymite trona-k</pre>	<pre>th(no3)4:5h2o th(so4)2 th2se3 thbr4 thenardite thf4 thi4 ths2 tiemannite t1 tobermorite-11a tobermorite-9a torbernite trevorite troilite tsumebite</pre>
713 715 717 719 721 723 725 727 729 731 733 735 737	<pre>th(oh)4 th2s3 th7s12 thc14 thermonatrite thf4:2.5h2o thorianite ti titanite tm tobermorite-14a todorokite tremolite tridymite</pre>	<pre>th(no3)4:5h2o th(so4)2 th2se3 thbr4 thenardite thf4 thi4 ths2 tiemannite tl tobermorite-11a tobermorite-9a torbernite trevorite troilite tsumebite u</pre>
713 715 717 719 721 725 727 729 731 733 735 737 739 741 743 745	<pre>th(oh)4 th2s3 th7s12 thc14 thermonatrite thf4:2.5h2o thorianite ti titanite tm tobermorite-14a todorokite tremolite tridymite trona-k tyuyamunite</pre>	<pre>th(no3)4:5h2o th(so4)2 th2se3 thbr4 thenardite thf4 thi4 ths2 tiemannite t1 tobermorite-11a tobermorite-9a torbernite trevorite troilite tsumebite</pre>
713 715 717 729 721 723 725 727 729 731 733 735 737 739 741 743 745 747	<pre>th(oh)4 th2s3 th7s12 thc14 thermonatrite thf4:2.5h2o thorianite ti titanite tm tobermorite-14a todorokite tremolite tridymite trona-k tyuyamunite u(co3)2 u(oh)2so4 u(so4)2</pre>	<pre>th(no3)4:5h2o th(so4)2 th2se3 thbr4 thenardite thf4 ths2 tiemannite tl tobermorite-11a tobermorite-9a torbernite trevorite troilite tsumebite u u(hpo4)2:4h2o u(so3)2 u(so4)2:4h2o</pre>
713 715 717 719 721 723 725 727 729 731 735 737 739 741 743 745 747 749	<pre>th(oh)4 th2s3 th7s12 thc14 thermonatrite thf4:2.5h2o thorianite ti titanite tm tobermorite-14a todorokite tremolite tridymite trona-k tyuyamunite u(co3)2 u(oh)2so4 u(so4)2 u(so4)2:8h2o</pre>	<pre>th(no3)4:5h2o th(so4)2 th2se3 thbr4 thenardite thf4 thi4 ths2 tiemannite tl tobermorite-11a tobermorite-9a torbernite trovilite troilite tsumebite u u(hpo4)2:4h2o u(so3)2 u(so4)2:4h2o u2c3</pre>
713 715 717 729 721 723 725 727 729 731 733 735 737 739 741 743 745 747 749 751	<pre>th(oh)4 th2s3 th7s12 thc14 thermonatrite thf4:2.5h2o thorianite ti titanite tm tobermorite-14a todorokite tremolite tridymite trona-k tyuyamunite u(co3)2 u(oh)2so4 u(so4)2 u(so4)2:8h2o u2f9</pre>	<pre>th(no3)4:5h2o th(so4)2 th2se3 thbr4 thenardite thf4 thi4 ths2 tiemannite t1 tobermorite-11a tobermorite-9a torbernite trevorite troilite tsumebite u u(hpo4)2:4h2o u(so3)2 u(so4)2:4h2o u2c3 u2c35</pre>
713 715 717 721 723 725 727 729 731 733 735 737 739 741 743 745 747 749 751 753	<pre>th(oh)4 th2s3 th7s12 thc14 thermonatrite thf4:2.5h2o thorianite ti titanite tm tobermorite-14a todorokite tremolite tridymite trona-k tyuyamunite u(co3)2 u(oh)2so4 u(so4)2:8h2o u2f9 u2o3f6</pre>	<pre>th(no3)4:5h2o th(so4)2 th2se3 thbr4 thenardite thf4 thi4 ths2 tiemannite tl tobermorite-11a tobermorite-9 torbernite trevorite trevorite troilite u u(hpo4)2:4h2o u(so3)2 u(so4)2:4h2o u2c3 u2o2c15 u2s3</pre>
713 715 717 729 721 723 727 729 731 735 737 739 741 743 745 747 749 753 755	<pre>th(oh)4 th2s3 th7s12 thc14 thermonatrite thf4:2.5h2o thorianite ti titanite ti todorokite tremolite tridymite trona-k tyuyamunite u(co3)2 u(so4)2 u(so4)2 u(so4)2 u(so4)2 u(so4)2 u(so36 u2se3</pre>	<pre>th(no3)4:5h2o th(so4)2 th2se3 thbr4 thenardite thf4 thi4 ths2 tiemannite tl tobermorite-11a tobermorite-9a torbernite trovilite troilite tsumebite u u(hpo4)2:4h2o u(so3)2 u(so4)2:4h2o u2c3 u2o2c15 u2c3 u3as4</pre>
$\begin{array}{c} 713\\ 715\\ 717\\ 719\\ 721\\ 725\\ 727\\ 729\\ 731\\ 733\\ 735\\ 737\\ 739\\ 741\\ 743\\ 745\\ 747\\ 749\\ 751\\ 755\\ 757\\ \end{array}$	<pre>th(oh)4 th2s3 th7s12 thc14 thermonatrite thf4:2.5h2o thorianite ti titanite tm tobermorite-14a todorokite tremolite tridymite trona-k tyuyamunite u(co3)2 u(oh)2so4 u(so4)2 u(so4)2:8h2o u2f9 u2o3f6 u2se3 u3o5f8</pre>	<pre>th(no3)4:5h2o th(so4)2 th2se3 thbr4 thenardite thf4 thi4 ths2 tiemannite t1 tobermorite-11a tobermorite-9a torbernite trevorite troilite tsumebite u u(hpo4)2:4h2o u(so3)2 u(so4)2:4h2o u2c3 u2c3 u3as4 u3p4</pre>
713 715 717 729 721 723 727 729 731 735 737 739 741 743 745 747 749 753 755	<pre>th(oh)4 th2s3 th7s12 thc14 thermonatrite thf4:2.5h2o thorianite ti titanite ti todorokite tremolite tridymite trona-k tyuyamunite u(co3)2 u(so4)2 u(so4)2 u(so4)2 u(so4)2 u(so4)2 u(so36 u2se3</pre>	<pre>th(no3)4:5h2o th(so4)2 th2se3 thbr4 thenardite thf4 thi4 ths2 tiemannite tl tobermorite-11a tobermorite-9a torbernite trovilite troilite tsumebite u u(hpo4)2:4h2o u(so3)2 u(so4)2:4h2o u2c3 u2o2c15 u2c3 u3as4</pre>
$\begin{array}{c} 713\\ 715\\ 717\\ 719\\ 721\\ 725\\ 727\\ 729\\ 731\\ 733\\ 737\\ 739\\ 741\\ 743\\ 745\\ 747\\ 753\\ 755\\ 757\\ 759\\ 757\\ 759\\ 763\\ 763\\ \end{array}$	<pre>th(oh)4 th2s3 th7s12 thc14 thermonatrite thf4:2.5h2o thorianite ti titanite tm tobermorite-14a todorokite tremolite tridymite trona-k tyuyamunite u(co3)2 u(oh)2so4 u(so4)2 u(so4)2:8h2o u2f9 u2o3f6 u2se3 u3o5f8 u3s5</pre>	<pre>th(no3)4:5h2o th(so4)2 th2se3 thbr4 thenardite thf4 thi4 ths2 tiemannite tl tobermorite-11a tobermorite-11a tobermite trevorite trevorite troilite tsumebite u u(hpo4)2:4h2o u(so3)2 u(so4)2:4h2o u2c3 u3as4 u3p4 u3se4</pre>
713 715 717 719 721 723 725 727 731 733 735 737 739 741 737 739 741 755 747 755 755 755 755 757 759 761 363 765	<pre>th(oh)4 th2s3 th7s12 thc14 thermonatrite thf4:2.5h20 thorianite ti titanite tm tobermorite-14a todorokite tremolite tridymite trona-k tyuyamunite u(co3)2 u(oh)2so4 u(so4)2 u(so4)2:8h20 u269 u203f6 u2se3 u3o5f8 u3s5 u3se5 u5o12c1 uas2</pre>	<pre>th(no3)4:5h2o th(so4)2 th2se3 thbr4 thenardite thf4 thi4 ths2 tiemannite tl tobermorite-11a tobermorite-11a tobermite trevorite trevorite troilite tsumebite u u(hpo4)2:4h2o u(so4)2:4h2o u2c3 u2c31 u3e4 u3p4 u3p4 u3se4 u4f17 uas ubr2c1</pre>
713 715 717 719 721 723 725 727 731 733 735 737 739 741 743 745 747 749 751 755 757 757 757 759 761 763 763	<pre>th(oh)4 th2s3 th7s12 thc14 thermonatrite thf4:2.5h2o thorianite ti titanite tm tobermorite-14a todorokite tremolite tridymite trona-k tryuyamunite u(co3)2 u(so4)2 u(so4)2 u(so4)2 u(so4)2 u(so4)2 u(so5f8 u3s5 u3s55 u3se5 u5o12c1 uas2 ubr2c12</pre>	<pre>th(no3)4:5h2o th(so4)2 th2se3 thbr4 thenardite thf4 thi4 ths2 tiemannite tl tobermorite-11a tobermorite-9a torbernite trevorite troilite tsumebite u u(hpo4)2:4h2o u(so3)2 u(so4)2:4h2o u2c3 u3as4 u3p4 u3se4 u4f17 uas ubr2c1 ubr3</pre>
713 715 717 719 721 723 725 727 731 733 735 737 739 741 737 739 741 755 747 755 755 755 755 757 759 761 363 765	<pre>th(oh)4 th2s3 th7s12 thc14 thermonatrite thf4:2.5h20 thorianite ti titanite tm tobermorite-14a todorokite tremolite tridymite trona-k tyuyamunite u(co3)2 u(oh)2so4 u(so4)2 u(so4)2:8h20 u269 u203f6 u2se3 u3o5f8 u3s5 u3se5 u5o12c1 uas2</pre>	<pre>th(no3)4:5h2o th(so4)2 th2se3 thbr4 thenardite thf4 thi4 ths2 tiemannite tl tobermorite-11a tobermorite-11a tobermorite trevorite troilite tsumebite u u(hpo4)2:4h2o u(so3)2 u(so4)2:4h2o u2c3 u2c3 u3e4 u3p4 u3p4 u3se4 u4f17 uas ubr2c1</pre>

773	ubrcl3	uc
775	ucl.94(alpha)	ucl2f2
777	ucl2i2	ucl3
779	ucl3f	ucl3i
781	ucl4	ucl5
783	ucl6	uclf3
785	ucli3	uf3
787	uf4	uf4:2.5h2o
789	uf5(alpha)	uf5(beta)
791	uf6	uh3(beta)
793	ui3	ui4
795	umangite	uo2(am)
797	uo2(aso3)2	uo2(io3)2
799	uo2(no3)2	uo2(no3)2:2h2o
801	uo2(no3)2:3h2o	uo2(no3)2:6h2o
803	uo2(no3)2:h2o	uo2(oh)2(beta)
805	uo2(po3)2	uo2.25
807	uo2.3333(beta)	uo2.6667
809	uo2br2	uo2br2:3h2o
811	uo2br2:h2o	uo2broh:2h2o
813	uo2cl	uo2cl2
815	uo2cl2:3h2o	uo2cl2:h2o
817	uo2cloh:2h2o	uo2f2
819	uo2f2:3h2o	uo2foh
821	uo2foh:2h2o	uo2foh:h2o
823	uo2hpo4	uo2hpo4:4h2o
825	u02s03	uo2so4
827	uo2so4:2.5h2o	uo2so4:3.5h2o
829		
	uo2so4:3h2o	uo2so4:h2o
831	uo3(alpha)	uo3(beta)
833	uo3(gamma)	uobr2
835	uobr3	uocl
837	uocl2	uocl3
839	uof2	uof2:h2o
841	uof4	uofoh
843	uofoh:.5h2o	up
845	up2	up2o7
847	up2o7:20h2o	upo5
849	uraninite	uranocircite
851	uranophane	us
853	us1.9	us2
855	us3	use
857	use2(alpha)	use2(beta)
859	use3	v
861	v2o4	v3o5
863	v4o7	vaesite
865	vivianite	W
867	wairakite	weeksite
869	whitlockite	wilkmanite
871	witherite	wollastonite
873	wurtzite	wustite
875	xonotlite	У
	yb	zincite
879	zircon	zn
	zn3(aso4)2	zncr2o4
883	znf2	znseo3:h2o
885	zoisite	zr

liquids

1 br2

quicksilver

* note - (eqpt/pcrsg) The pure liquids block has not been written on the datal and datalf files, because the EQ3NR and EQ6 codes presently do not treat non-aqeuous liquids.

gases

1 3 7 9 11 13 15 17 23 25 27 29 31 33 5 35	ag(g) ar(g) bf3(g) c(g) cd(g) cl2(g) co2(g) h2s(g) hc1(g) hf(g) hi(g) hi(g) li(g) li(g) n2(g) ne(g) o2(g)	al(g) b(g) br2(g) ca(g) ch4(g) co(g) f2(g) h2o(g) hbr(g) hbr(g) hg(g) i2(g) kr(g) mg(g) na(g) nh3(g) pb(g)
33	ne(g)	nh3(g)
35 37	o2(g) rb(q)	pb(g) rn(q)
39	s2(g)	si(g)
41	sif4(g)	sn(g)
43	so2(g)	tc2o7(g)

55 57 61 63 65 67 69 71	u2cl10(g) u2f10(g) ubr2(g) ubr4(g) ucl2(g) ucl4(g) ucl6(g) uf2(g) uf4(g) uf4(g) ui2(g) ui2(g) ui2(g) ui2(g)	<pre>ti(g) u(g) u2cl8(g) ubr(g) ubr3(g) ucl(g) ucl3(g) ucl5(g) uf5(g) uf5(g) ui(g) ui3(g) ui3(g) uo2cl2(g) uof4(g) zn(g)</pre>
solid	solutions	
1	biotite	
	annite	phlogopite
2	carbonate-calcite calcite	magnagita
	rhodochrosite	magnesite siderite
	smithsonite	strontianite
3	chlorite-ss	Scioncianice
5	clinochlore-14a	daphnite-14a
4	clinoptilolite-ss	<u>F</u>
	clinoptilolite-ca	clinoptilolite-cs
	clinoptilolite-k	clinoptilolite-na
	clinoptilolite-nh4	clinoptilolite-sr
5	epidote-ss	
	clinozoisite	epidote
б	garnet-ss	
-	andradite	grossular
7	olivine	forsterite
8	fayalite orthopyroxene	Iorsterite
0	enstatite	ferrosilite
9	plagioclase	TETTODITICE
-	albite high	anorthite
	sanidine high	
10	sanidine-ss	
	albite high	sanidine high
	anorthite	
11	saponite-tri	
	saponite-ca	saponite-h
	saponite-k	saponite-mg
12	saponite-na smectite-di	
12	beidellite-ca	beidellite-k
	beidellite-ma	beidellite-na

beidellite-k beidellite-na montmor-k montmor-na nontronite-k nontronite-na

The slist file for the data0.hmw.R10 data file:

EQPT Species List File:

beidellite-mg montmor-ca

nontronite-ca nontronite-mg

montmor-mg

no. of elements on the data file = 9
 the dimensioned limit = 100
no. of aqueous species in the master set = 13
 the dimensioned limit = 500

element	=	0	,	atwt	=	15.99940
element	=	ca	,	atwt	=	40.07800
element	=	cl	,	atwt	=	35.45270
element	=	h	,	atwt	=	1.00794
element	=	С	,	atwt	=	12.01100
element	=	k	,	atwt	=	39.09830
element	=	mg	,	atwt	=	24.30500
element	=	na	,	atwt	=	22.98977
element	=	s	,	atwt	=	32.06600

aqueous

1 3 7 9 11 13 15 17	h2o cl- hco3- mg++ so4 co2(aq) oh- hso4- mgoh+	ca++ h+ k+ na+ o2(g) co3 caco3(aq) mgco3(aq)
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minerals

7 9 11 13 15 17 19 21 23 25 27 29 31 33	<pre>burkeite ca4cl2(oh)6:13h2o calcite dolomite gaylussite gypsum hexahydrite k3h(so4)2 kainite kieserite leonite mercallite misenite</pre>	antarcticite aragonite bischofite brucite ca2cl2(oh)2:h2o cacl2:4h2o carnallite epsomite glauberite halite k2co3:3/2h2o k8h4(co3)6:3h2o k8h4(co3)6:3h2o kalicinite knaco3:6h2o magnesite mirabilite na2co3:7h2o
27	kieserite	knaco3:6h2o
	na3h(so4)2	na4ca(so4)3:2h2o
37		natron
39 41	nesquehonite picromerite	oxychloride-mg pirssonite
43		portlandite
45		syngenite
47		thenardite
	thermonatrite	trona
51	trona-k	

liquids

0 none

* note - (eqpt/pcrsg) The pure liquids block has not been written on the datal and data1f files, because the EQ3NR and EQ6 codes presently do not treat non-aqeuous liquids.

gases

1 co2(g) 3 o2(g)

h2(g)

solid solutions

0 none