

General Thermodynamic Package (GTP)

Basic description and documentation, version 3.21

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This is a preliminary description of the data structure and subroutines in the thermodynamic model package GTP. A background to this initiative to develop a free thermodynamic software can be found in[1].

This documentation is neither complete nor up to date with the most recent version. The whole structure (and depending subroutines) will be revised when all major parts have been implemented and tested.

The OC software is written in the new Fortran standard and the main reason to use Fortran is that it is specially designed for numerical calculations and there are many numerical libraries available. But it is also due to the age of the developer. Fortran is useful for parallel processing and thermodynamic calculations are needed in many simulation software for materials processee. Extensive use is made of the datatyping available in the new Fortran.

Documentation updating software

One very difficult thing with software that is alive and changing is to update the documentation. So my feeling for documentation is that once a software is documented it is dead as it is almost impossible to update source code and documentation in parallel.

In an attempt to challenge this feeling I have written a simple documentation update software that uses an existing documentation text written as a LaTeX file together with the source code to compare critical parts in the source code as “verbatim” parts. In the documentation there are text with additional explanations of the verbatim parts and each verbatim section is in a separate LaTeX (sub)section. The documentation software compares the verbatim sections in the existing documentation with those in the new source code and whenever it finds a difference the developer is notified. If there are changes within an existing verbatim section the old and new verbatim parts are both written in the new documentation and the developer must update the text around these verbatim parts. If there are new verbatim sections in the source code the developer must edit the old documentation to add a (sub)section for the new verbatim and rerun the documentation software. A verbatim section which has changed place in the source code also requires editing of the existing documentation. This is a safe method to find what has changed in the source as it is a quite task for the developer to add new verbatim sections in the source code. But running the documentation several times with editing the LaTeX file can be quite tedious.

The verbatim sections in the source code is enclosed by

```
!\begin{verbatim}  
  
critical part of the software for example a function declaration  
  
!\end{verbatim}
```

The “begin” and “end” must be left justified on the line.

The verbatim parts can be inserted anywhere in the source code, typically they enclose important sections like data structure definitions and declarations of subroutines and functions. The documentation software detect new or changed verbatim sections as well as verbatim sections that has disappeared and writes a new LaTeX documentation file which require editing to describe the changes and to add crossreferences. Using this software I hope to keep the software documented also during the development stage. This document is an example of using this software.

The subroutines are documented in the order they appear in the source code. That order is not always logical and sometimes a major restructuring will be made. A table with the subroutines can be generated in the LaTeX file by adding a line

```
!\addtotable <name of function or subroutine>
```

at each subroutine.

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1 Introduction

In a model for a thermodynamic system one has elements, species (combination of elements with fixed stoichiometric ratios and a possible charge) and phases. The thermodynamic properties of the system can be described by the Gibbs energy for each phase and the Gibbs energy depend on the temperature, T , pressure, P and the amounts of the constituents of the phase, the constituents are a subset of the species.

At equilibrium at constant T , P and composition the Gibbs energy of a system is at a minimum. With suitable transformations the Gibbs energy can be used to find the equilibrium of a system for all other sets of conditions (except when a phase has a miscibility gap in the volume). From the Gibbs energy one can calculate many important properties of the system like the set of stable phases and their constitution, the chemical potentials of the components, the heat content, heat capacity, volume, thermal expansion and much more. It is also possible to calculate phase diagrams and the model functions can be used to extrapolate to non-equilibrium states. Some thought has been spent on modeling special physical properties like ferro-magnetism using additional internal variables like the Curie Temperature and how elastic constants (which are not constant) can be included in the modeling.

The method used for storing composition dependent properties of the Gibbs energy parameters is suitable also for other properties that depend on the phase like mobilities, resistivities, viscosities etc and there are provisions to include such data in the software.

The use of the GTP model package to calculate equilibria in multicomponent systems, for calculation of property and phase diagrams and for use in other application oriented software is explained in separate documents and papers.

1.1 New features in current and previous versions

Version 5 of OC was made available early 2018 at the OC homepage[2] but there was no update of the documentation. Frequent updates of the code has been made at github[3], currently (July 2019) version 5.045. But this is the first update of this documentation. I hope to be able to update the other parts and the User Guide later this year.

OC is shaping up fairly well and there are a few serious users, it is interesting to note that there are several interfaces to OC available on github, even for Python. But OC is now so large it is difficult even for me to find the correct place to update the code and often changes has to be made in many different files. This means documentation becomes even more important! I hope to release version 6 of OC together with a complete documentation update early 2020.

There has been quite a lot of changes in the model package. Recently the UNIQUAC model for fluid was added and earlier, see section 8.15.12 and also the Einstein model, section 2.3.6, for heat capacity and the two-state model for amorphous/liquid models, section 2.3.7. A model for extrapolation of solids above their melting T has also been added (based on the

equi-entropy temperature, EET) but that is part of the minimizing package as it is related to the equilibrium calculation, not a part of the description of a single phase. However, the EET criterion is activated by setting a positive value in the global datastructure described in section 5.1.

It is now possible to save all workspaces, except results from STEP and MAP commands on an unformatted Fortran file. This makes it possible to save the current state of a calculation on a file and later retrieve all the data from this file. This is currently very useful for simulations and for assessments and all experimental data and optimizing coefficients can be saved.

Version 4 of OC was released early 2017. A prerelease is available at the opencalphad repository at github[3]. There has been several improvement in various parts. The numeric routines now use LAPACK and BLAS and if a user has a tailored version of these for his computer there may a gain in speed about 5-10%. To simplify the installation there is a subset of the necessary LAPACK and BLAS routines included in this distribution and even these improved the performance with about 20% compared to the previous numeric routines. Other features added are the possibility to use P as variable and V as condition and expressions using amount of moles of components. It is also possible to define other components than the elements and one can calculate and plot isothermal phase diagrams.

There is no new thermodynamic models implemented. For the equilibrium calculations a paper has been published[4], for more details see the documentation of the equilibrium module, For mapping some improvements have been in the step control but the mapping is very sensitive to the start point. There has been some minor changes in the graphics interface to GNUPLOT. There has been great interest using OC for application software and another paper was published recently[5] together with several users, there will hopefully be a more extensive documentation of the OC Application Software Interface (OCASI) with this release. The possibility to use OC with C++ has been greatly simplified with the OC-isoC interface.

The possibility to save and read the whole datastructure for an equilibrium, mapping or assessment on a file has not been possible to achieve. That will be next release.

The release of version 3 of OC is available at the opencalphad webpage[2] since February 2016. It included several improvements in the already existing software but most important a new assessment procedure to fit model parameters to experimental and theoretical data. This has added a few new data structures and commands to the GTP package. Version 3 can also be used for parallel calculations of several equilibria using the OpenMP package. A significant increase in speed has been noted.

Version 2 of OC was released in February 2015 and the main new feature of this version was the step and map procedures which are documented separately. It also included derivatives of state variables, “dot derivatives”, and some improvements of the minimization procedure and main change in the GTP module was implementing the ionic liquid model. This did not really change much in this documentation, the code for handling the variable ratio of

sublattices was been integrated in the existing subroutines.

1.2 Features to be added in future versions

A software is never finished but each version has to be tested and debugged before new features can be added. It is now possible to save/read unformatted data on a file but not yet results from STEP and MAP.

There are a still models I want to add to handle SRO. As OC is a free software it is possible for any interested scientist to contribute, it is one of the reasons I am writing this documentation.

1.3 The Gibbs energy function

The model package provides the possibility to calculate the Gibbs energy, G , for many phases for any value of T, P and its constitution using the models implemented. In addition it can calculate the first and second derivatives of G with respect to T, P and the constitution. These values can be used directly or within a minimization package developed in parallel, to handle equilibrium calculations in multicomponent system with several phases for various external conditions. The basic formula for the Gibbs energy of a phase α , independent of the model, is taken from the book by Lukas, Fries and Sundman[6].

$${}^{\alpha}G_M^{\alpha} = {}^{\text{srf}}G_M^{\alpha} - T {}^{\text{cfg}}S_M^{\alpha} + {}^E G_M^{\alpha} + {}^{\text{phys}}G_M^{\alpha} \quad (1)$$

where ${}^{\text{srf}}G_M^{\alpha}$ is the surface of reference containing parameters for the endmembers, ${}^{\text{cfg}}S_M^{\alpha}$ is the configurational entropy, ${}^E G_M^{\alpha}$ is the excess Gibbs energy which has interaction parameters within the phase and ${}^{\text{phys}}G_M^{\alpha}$ is the contribution to the Gibbs energy due to some physical properties, like ferromagnetism, that are modeled separately.

The Gibbs energy is modeled for a formula unit of the phase, the formula unit defined by the crystallography, and the amount of the components for a formula unit of a phase can vary because the constituents can be for example molecules in a gas phase or vacancies occupying vacant lattice sites. The composition dependence of the modeled Gibbs energy depend on the constituent fractions, denoted y_i , possibly with a sublattice index and the mole fractions of a component A in the phase α can be calculated using the formula:

$$x_A^{\alpha} = \frac{\sum_s a_s^{\alpha} \sum_j b_{Aj} y_{js}^{\alpha}}{\sum_s a_s^{\alpha} \sum_B \sum_j b_{Bj} y_{js}^{\alpha}} \quad (2)$$

where a_s^{α} is the number of sites on sublattice s , b_{Aj} is the stoichiometric ratio of element A in species j and y_{js}^{α} is the constituent fraction of species j on sublattice s . The summation

over B is for all components. The sum of site fractions in a sublattice s is unity:

$$\sum_i y_{is} = 1 \quad (3)$$

The terms and factors used in this equation are explained below and for further details please read the book by Lukas et al.

1.4 Additions from physical properties

For many phases there are additions from different physical properties, $^{\text{phys}}G_M$. In most cases these are defined per mole atoms whereas the Gibbs energy in eq. 1 is defined per mole formula unit. It means we have:

$$^{\text{phys}}G_M^\alpha = N^\alpha \cdot ^{\text{phys}}G_m^\alpha \quad (4)$$

where N^α is the number of moles of atoms in α and $^{\text{phys}}G_m^\alpha$ the expression for the Gibbs energy of the addition per mole atom. See also sections 2.3.5, 3.2.2 and 5.6.9

2 Basis

The basic terms and concepts of the GTP model package are explained here. Depending on your background some definitions may not agree with what you are used.

2.1 Elements

The elements have a sequential index arranged alphabetically. They have a one or two-letter symbol and a mass. The symbol must be letters A-Z all in upper case. In addition a "name" of the element is stored, like "iron" for Fe, the name of a reference phase, the value of H298-H0 for the reference phase and S298 for the reference phase. The element symbol must be unique.

OC is case insensitive, it means that an element written as Fe is the same as fe or FE or fE. The element Cobalt can be written co and to specify the species carbon monoxide we must write clo or clo1 if there is also clo2 in the system.

The electron and vacancy are entered as element -1 and 0 respectively but they are not really treated as elements except that they can be included in species. The electron has the symbol /- but one can also use /+ to denote a positive charge (or /-1). Vacancies are denoted "Va" and have no fixed amount and will always have zero chemical potential at equilibrium. They will thus not affect the Gibbs phase rule.

An element can be suspended, that means it can be hidden from application programs. If an element is suspended then all species that include this element are also suspended (implicitly) and also any phases which cannot exist unless the species is present as constituent, i.e. the species is the only constituent in a sublattice.

2.2 Species

The species are stoichiometric combinations of elements. They have a symbol and a stoichiometric formula. The elements, including the vacancy, are the simplest species. The stoichiometry can be non-integer like a species $\text{AlO}_{1.5}$. The symbol must start with a letter A-Z and can contain letters (case insensitive), digits, the period ".", the underscore character "_", the slash "/", the minus sign "-" and the plus sign "+". No other special characters are allowed, for example parenthesis are not allowed.

The symbol of the species is often the same as the stoichiometric formula but it is not necessary. For example the molecule $\text{C}_2\text{H}_2\text{Cl}_2$ exists in 2 configurations either with Cl on the same side or opposite. These can be identified by giving these the names `C2H2CL2_CIS` and `C2H2CL2_TRANS`.

In addition the mass is stored but this is redundant as it is calculated from the element masses. The species can have a charge like $\text{Al}/+3$ or $\text{O}/-2$. The charge can be non-integer.

Note that a species does not represent a phase, just a stoichiometric formula. As constituents of a phase the species are associated with that phase. The species H_2O can be a constituent of ice, liquid water and gas. Thus species have no thermodynamic data. Their only function is to be constituents of phases where they can be assigned data. A species must not be confused with a phase just because many phases have a fixed stoichiometric formula. In this package the same species can be a constituent of many phases, also phases with variable composition.

When entering the stoichiometric formula for a species one may need to use stoichiometric numbers to separate one letter elements as `OC` is case insensitive. For example to distinguish CO as cobalt from C1O1 as carbon monoxide the digit 1 can be used after the element letter C. But otherwise the stoichiometric number 1 is not needed if the element has a two letter symbol, for example `CAO` can be used for a species with one atom of CA and one atom of O. One cannot use parenthesis in species symbols or when entering the stoichiometry, the stoichiometry of $\text{Ca}(\text{OH})_2$ must be entered as `CAO2H2` (or `O1CAO1H2` or any other way giving the same stoichiometry).

The fact that a species has a symbol which is independent of its stoichiometry can be used to simplify complicated chemical stoichiometries.

It is important to keep in mind that a species is identified with its symbol, not its stoichiometry. Thus a species with the symbol `c1o1` cannot be found by giving the symbol `o1c1`.

2.3 Phases

All thermodynamic data that are used in calculations are stored as part of a phase. The Gibbs energy of the phase is modeled as a function of temperature (T), pressure (P) and the constitution of the phase (Y) as given by eq. 1. Many different thermodynamic models can be used and each phase can have a unique model and also several different “additions” to this model, like magnetic and pressure models. But all phases are accessed from outside the model package in the same way. Thus the minimizer which is used to find the equilibrium state of a system does not depend on any particular model feature.

From outside the model package one can obtain information of the constitution of the phase, i.e. which constituents and their fractions (Y). The constituents are species (with fixed stoichiometry). As the compound energy formalism (CEF) is a very general and flexible model framework this has been the first implemented. In version 2 of OC the ionic liquid model is also implemented. There is no special documentation of that model in this text as it is described in the book by Lukas, Fries and Sundman [6].

In CEF one can define sublattices with specific species (elements or molecules) as constituents and the constituent fractions on each sublattice must be unity. The constituents are assumed to mix randomly on each sublattice. For a general description of CEF please consult Lukas et al.[6].

One may implement models that do not use sublattices and have non-random entropy in GTP, for example CVM or quasichemical models. Inside the CEF framework this can be done by treating the phase as having a single lattice and the probabilities of the different clusters are fractions and the clusters are species. Inside the model package the appropriate non-random configurational entropy is then evaluated. The reason to base the model package on the constitution and not the composition of the component is to avoid a two-level minimization when calculating the equilibrium. But there is nothing, except speed, preventing from using the mole fractions as external variables for some models and implement an internal minimization to determine the configuration which will give the minimum Gibbs energy for the given external conditions.

There are some models (not implemented in OC) which allow a variable stoichiometry of the constituents of the phase. Such models must be implemented as a two-level minimization where the model from outside the model package depend only the mole fractions of the elements or some other set of components. It may also be necessary to have some special way of entering parameters for such a model.

2.3.1 Phase specification

A phase has a name and some model information. If it is a CEF model it has number of sublattices, the sites in each sublattice (a fixed real value) and a list of constituents on each sublattice. One may have a single constituent in a sublattice. An empty sublattice can have the vacancy as the only constituent.

The term "endmember" refers to a combination of one constituent in each sublattice. This defines a compound and this is the origin of the name "compound energy formalism". When writing an endmember the constituents are separated by a colon.

The thermodynamic model parameters are entered for the endmembers and possible interaction parameters.

The ionic liquid model has variable number of sites for cations and anions to maintain electro neutrality. This must be taken care of when calculating the mass balance during with the minimization.

When entering an endmember parameter the software writes the amount and reference states of the elements this is referred to. Care must be taken that the parameter value correspond to the correct amount of atoms as this can vary between endmembers of the same phase. For example if the FCC phase is defined with 2 sublattices with one site in each, the endmember G(FCC,TI:VA) is for one mole of atoms whereas G(FCC,TI:C) is for two moles of atoms.

2.3.2 Interaction parameters

For interaction parameters one must add one, two or three additional constituents in any sublattice separated by a comma from the other constituents in the same sublattice. The fractions of the constituents specified in the parameter should be multiplied with the parameter value.

An interaction parameter may also have a degree specifying that it has a more complex composition dependence. The degree is a digit 0 to 9 given after a semicolon. The default degree is zero. The meaning of the degree depend on the model, for a binary interaction in CEF it is by default the power in a Redlich-Kister (RK) series:

$$L_{ij} = \sum_{\nu=0} (y_i - y_j)^{\nu} L_{ij}^{\nu} \quad (5)$$

where y_i and y_j are the constituent fractions on the same sublattice and ν is the degree of the parameter. For $\nu = 0$ the parameter is composition independent.

Each such RK coefficient is a function of temperature and pressure. There can also be different extrapolation models like Muggianu, Kohler and Toop but they are not implemented yet.

For a ternary parameter there is also possibilities to have composition dependence:

$$L_{ijk} = v_i^0 L_{ijk} + v_j^1 L_{ijk} + v_k^2 L_{ijk} \quad (6)$$

$$v_i = (y_i + (1 - y_i - y_j - y_k))/3 \quad (7)$$

$$v_j = (y_j + (1 - y_i - y_j - y_k))/3 \quad (8)$$

$$v_k = (y_k + (1 - y_i - y_j - y_k))/3 \quad (9)$$

where y_i, y_j and y_k are the constituent fractions on the same sublattice. The v_i are the same as y_i in the ternary system and the additional terms have been added to make this contribution symmetrical in higher order systems.

2.3.3 Model parameters identifiers

In addition to calculate the Gibbs energy for a phase the user can also enter parameters for other properties. These parameters can depend on T, P and the constitution of the phase. In this package all parameters are identified by

- a property symbol, listed in Table 1
- the phase name,
- the constituent array (specifying the constituents in each sublattice, the fraction of which are multiplied with the parameter, see also section 2.3.4), and
- a degree with a value 0 to 9, usually a Redlich-Kister power, see section 2.3.2.

Some examples are G(FCC,FE), G(SIGMA,FE:CR:CR), G(LAVES,FE,MO:MO;2), TC(BCC,FE) or MQ&FE(FCC,FE), see also 2.3.8.

The property symbol G is used for the Gibbs energy. Other properties symbols like TC is used for the Curie temperature and MQ&XY for the mobility of the element XY the the phase.

This means it is possible to use models that does not give any contribution to the Gibbs energy, like the mobility, and let the gtp model package to handle the T, P and composition dependence of the model parameters like resistivity, viscosity etc. A tentative list of properties that have been defined is given in Table 1.

In case the parameters should affect the Gibbs energy, like the Curie temperature, an “addition” subroutine must be written using the value of this parameter. See section 5.6.9.

The TC and BMAG parameters are included for compatibility with the current magnetic model, most of the others are just tentative.

2.3.4 Wildcards in model parameters

For phases with several sublattices a model parameter may be independent of the specific constituent in a sublattice. This can be indicated by used an asterix, “*”, instead of a constituent in that sublattice. This feature can also be used to use model parameters that represent “bonds” rather than endmember energies. For a parameter with a wildcard in a sublattice the fraction for that sublattice is replaced by the factor 1.0.

Table 1: Model parameter identifiers. Note a model identifier can have a suffix D indicating it belongs to the disordered fraction set, see section 2.5

Index	Symbol	T	P	Specification	Meaning
1	G	T	P		Energy
2	TC	-	P		Combined Curie/Neel T
3	BMAG	-	-		Average Bohr magneton numb
4	CTA	-	P		Curie temperature
5	NTA	-	P		Neel temperature
6	IBM	-	P	<constituent#sublattice>	Individual Bohr magneton num
7	THET	-	P		Debye or Einstein temp
8	V0	-	-		Volume at T0, P0
9	VA	T	-		Thermal expansion
10	VB	T	P		Bulk modulus
11	G2	T	P		Liquid two state parameter
12	LAMB	-	P		Thermal conductivity
13	MQ	T	P	<constituent#sublattice>	Mobility activation energy
14	MF	T	P	<constituent#sublattice>	RT*ln(mobility freq.fact.)
15	MG	T	P	<constituent#sublattice>	Magnetic mobility factor
16	THT2	-	P		Smooth step function T
17	DCP2	-	P		Smooth step function value
18	VISC	T	P		Viscosity
19	LPX	T	P		Lattice param X axis
20	LPY	T	P		Lattice param Y axis
21	LPZ	T	P		Lattice param Z axis
22	LPTH	T	P		Lattice angle TH
23	EC11	T	P		Elastic const C11
24	EC12	T	P		Elastic const C12
25	EC44	T	P		Elastic const C44
26	UQT	T	P	<constituent#sublattice>	UNIQUAC residual parameter
27	RHO	T	P		Electric resistivity
28	HMVA	T	P		Enthalpy of vacancy form.
29	TSCH	-	P		Schottky anomaly T
30	CSCH	-	P		Schottky anomaly Cp/R.
31	NONE	T	P		Unused

2.3.5 Physical models

Traditionally CALPHAD models describes the variation of the Gibbs energy as a function of T , P and composition without bothering about the origin of the contributions. In some cases this is no so good and for the contribution due to ferromagnetic transitions one has introduced a more physical contribution that depend on two new variables, the Curie temperature and the Bohr magneton number, both of which depend on the constitution. Such *additions*

to the traditional Gibbs energy expression may become more frequent when also phonon contributions are modeled down to 0 K as a function of a constitution dependent Debye temperature. Hopefully this can lead to better extrapolations to higher temperatures as well as multicomponent systems.

There are no description of the well established additions here because they can be found in Lukas et al [6]. But some of the new models will be described here.

The Gibbs energy in eq. 1 is per mole formula unit of the phase and so is also the $^{\text{phys}}G_M$ term but the expressions found in textbooks are always per mole of atom. Thus the Gibbs energy expression for the physical model must be multiplied with the number of atoms per formula unit of the phase. This is done by setting a bit ADDPERMOL in the addition record, see section 3.2.2. How to do that will be described in (a future update of) the User Guide.

2.3.6 Einstein heat capacity model

The Einstein model is not in Lukas [6] and its Gibbs energy contribution, obtained by integrating the heat capacity contribution due to the Einstein model is:

$$\Delta G_M^{\text{Ein}} = \frac{3}{2}R\theta + 3RT \ln(1 - \exp(-\frac{\theta}{T})) \quad (10)$$

where θ is the Einstein temperature which can vary with composition and P but not T . The Einstein heat capacity model is preferred to the Debye model because it can be integrated to a closed expression and both models must anyway have additional terms to fit the heat capacity data for any real element. The Einstein T, θ , is not a materials constant but a value fitted to the low T heat capacity curve for the elements. In some cases different θ can be used in different T regions.

The composition dependence of θ is best described by adding $\ln(^{\circ}\theta_{E,i})$ for each constituents i (including possibly some excess parameters) and then calculate:

$$\theta = \exp(\sum_i y_i \ln(^{\circ}\theta_{E,i})) \quad (11)$$

$$\frac{\partial \theta}{\partial y_i} = \theta \frac{\partial}{\partial y_i} (\sum_j y_j \ln(^{\circ}\theta_{E,j})) \quad (12)$$

We have to derive the first and second derivatives both wrt T and composition as θ will depend on the composition.

$$\begin{aligned} G_M &= \frac{3}{2}R\theta + 3RT \ln(1 - \exp(-\frac{\theta}{T})) \\ \frac{\partial G_M}{\partial T} &= 3R \ln(1 - \exp(-\frac{\theta}{T})) + 3RT \frac{-\frac{\theta}{T^2} \exp(-\frac{\theta}{T})}{1 - \exp(-\frac{\theta}{T})} \end{aligned} \quad (13)$$

$$= 3R \ln(1 - \exp(-\frac{\theta}{T})) - \frac{3R\frac{\theta}{T}}{\exp(\frac{\theta}{T}) - 1} \quad (14)$$

$$\frac{\partial G_M}{\partial y_i} = R(\frac{3}{2} + \frac{2}{\exp(\frac{\theta}{T}) - 1}) \frac{\partial \theta}{\partial y_i} \quad (15)$$

$$\begin{aligned} \frac{\partial^2 G_M}{\partial T^2} &= 3R(-\frac{\frac{\theta}{T^2}}{\exp(\frac{\theta}{T}) - 1} - \frac{-\frac{\theta}{T^2}(\exp(\frac{\theta}{T}) - 1) + (\frac{\theta}{T})^2 \exp(\frac{\theta}{T})}{(\exp(\frac{\theta}{T}) - 1)^2}) \\ &= 3R(-\frac{\frac{\theta}{T^2}}{\exp(\frac{\theta}{T}) - 1} + \frac{\frac{\theta}{T^2}}{\exp(\frac{\theta}{T}) - 1} - \frac{\theta^2}{T^3} \frac{\exp(\frac{\theta}{T})}{(\exp(\frac{\theta}{T}) - 1)^2}) \\ &= -3R \frac{\theta_E^2}{T^3} \frac{\exp(\frac{\theta}{T})}{(\exp(\frac{\theta}{T}) - 1)^2} \end{aligned} \quad (16)$$

The heat capacity at constant P is:

$$C_P = -T \frac{\partial^2 G_M}{\partial T^2} = 3R(\frac{\theta}{T})^2 \frac{\exp(\frac{\theta}{T})}{(\exp(\frac{\theta}{T}) - 1)^2} \quad (17)$$

which is the correct expression for C_V according to Einstein:

$$C_V = 3R(\frac{\theta}{T})^2 \frac{\exp(\frac{\theta}{T})}{(\exp(\frac{\theta}{T}) - 1)^2} \quad (18)$$

This gives $C_P = 0$ at 0 K. The value for S is also zero at 0 K from eq. 14 as $\exp(-\frac{\theta}{T}) = 0$ when $T = 0$ as $\theta > 0$.

$$S(T = 0) = -\left(\frac{\partial G}{\partial T}\right)_{T=0} = 0 \quad (19)$$

and for the Gibbs energy the value at $T = 0$ will be $\frac{3}{2}R\theta$.

2.3.7 The liquid twostate model

The liquid two-state model is not in Lukas [6] and the model equation and some partial derivatives are given here as part of the documentation.

In the liquid two-state model the Gibbs energy of the liquid is described as:

$$G_M^{\text{am+liquid}} = G_M^{\text{Ein}}(\theta) - G_M^{\text{2state}} + G_M^{\text{corr}} \quad (20)$$

$$G_M^{\text{2state}} = RT \ln(1 + \exp(-\frac{G_2}{RT})) \quad (21)$$

where G_M^{Ein} is the Gibbs energy of an amorphous state with an Einstein heat capacity contribution which depends on an estimated composition dependent θ for the amorphous state. The function G_M^{2state} describes the transition from the amorphous to the liquid state. At

0 K it gives zero contribution to the Gibbs energy, entropy and heat capacity and above the melting T it should describe the real liquid properties. The parameter $G2$ can depend on T, P and the composition of the liquid. For a substitutional liquid that would be

$$G2 = \sum_i (x_i {}^\circ G2_i + (\sum_{j>i} x_j G2_{ij} + \dots)) \quad (22)$$

where x_i is the mole fraction of component i . For a pure element ${}^\circ G2_i$ is:

$${}^\circ G2_i = a0_i + a1_i T + a2_i T \ln(T) + \dots \quad (23)$$

where $a0_i$ is related to the enthalpy of melting and $a1_i$ to the entropy of melting (close to R for many metals according to Richard's rule). But $a0_i, a1_i$ and $a2_i$ and possibly more terms can be treated as parameters to be fitted to all the experimental data like the melting T of the crystalline solid, the enthalpy and entropy of melting and the heat capacity of the stable liquid.

The final term G_M^{corr} in eq. 20 can be used to adjust the Gibbs energy of the stable liquid in addition to the previous terms. In fact this G_M^{corr} is the parameter used to describe the liquid phase without the two-state model. It is also composition dependent and for a substitutional liquid it can be written:

$$G_M^{\text{corr}} = \sum_i (x_i {}^\circ G_i^{\text{corr}} + (\sum_{j>i} x_j G_{ij}^{\text{corr}} + \dots)) \quad (24)$$

For a pure element i :

$${}^\circ G_i^{\text{corr}} = b0_i + b1_i T + b2_i T^2 \quad (25)$$

where $b0_i$ can be interpreted as the enthalpy difference between a metastable liquid and a (slightly less metastable) amorphous phase at 0 K for component i . $b1_i$ is an entropy at $T = 0$ K due to its disordered state and $b2_i$ can be used to fit the properties of the amorphous phase. One can use higher powers in T but not any $T \ln(T)$ or negative powers because such terms would give a contribution to the heat capacity at 0 K.

At 0 K eq. 20 should represent the Gibbs energy of the amorphous state with $C_P = 0$. There can be a finite entropy at 0 K because the amorphous phase is not a defect free crystal. At increasing T the heat capacity for the amorphous state should follow an Einstein curve.

Anyway, to implement this model we have to calculate some derivatives of $G_M^{2\text{state}}$ with respect to T and constitution, x_i of the liquid. We must also use the derivatives of the Einstein heat capacity function for the amorphous phase in section 2.3.6. For simplicity the superscript "2state" is skipped in the following equations. Note that $G2$ and all its derivatives are calculated together with other parameters for each iteration.

$$G_M = RT \ln(1 + \exp(-\frac{G2}{RT})) \quad (26)$$

$$\begin{aligned}
\frac{\partial G_M}{\partial T} &= R \ln(1 + \exp(-\frac{G2}{RT})) + RT \frac{(-\frac{1}{RT} \frac{\partial G2}{\partial T} + \frac{G2}{RT^2}) \exp(-\frac{G2}{RT})}{1 + \exp(-\frac{G2}{RT})} \\
&= R \ln(1 + \exp(-\frac{G2}{RT})) + \frac{-\frac{\partial G2}{\partial T} + \frac{G2}{T}}{\exp(\frac{G2}{RT}) + 1}
\end{aligned} \tag{27}$$

$$\frac{\partial G_M}{\partial y_i} = -\frac{\partial G2}{\partial y_i} \frac{1}{\exp(\frac{G2}{RT}) + 1} \tag{28}$$

And for the second derivatives:

$$\frac{\partial^2 G}{\partial T^2} = (-\frac{\partial^2 G2}{\partial T^2} + \frac{(\frac{G2}{T})^2 + (\frac{\partial G2}{\partial T})^2 - 2\frac{G}{T} \frac{\partial G2}{\partial T}}{RT(1 + \exp(-\frac{G2}{RT}))}) / (1 + \exp(\frac{G2}{RT})) \tag{29}$$

For the moment I skip the 2nd derivatives wrt fractions ...

2.3.8 Other properties than the Gibbs energy

The Gibbs energy parameters has property index 1 (one). There are a number of predefined properties in the code, that is the Curie temperature (index 2), average Bohr magneton number (3) etc. These are defined in the subroutine `init.gtp` and a tentative list of predefined identifiers is given in Table 1. More types can be added but one must choose a unique symbol for each and each is assigned a sequential index. The symbol must not be mistaken for a state variable symbol either. In the property records, see 5.6.6, for endmembers and interactions this property index is used to identify the property.

The mobility of an element in a phase depends on the constitution and the temperature but the mobility does not contribute to the Gibbs energy. Anyway it is possible to define and enter mobilities in GTP to handle their composition dependence. The values of these, and any other separate property like the Curie temperature, can be obtained anytime by a special subroutine call.

One must consider that the mobility for each constituent depend individually on the constitution and a symbol `MQ&X(BCC)` is used to define the mobility of X in the BCC phase as a function of its constitution. Thus `MQ&FE(BCC,FE)` is the self diffusion of FE in BCC and `MQ&FE(BCC,Cr)` is the mobility of FE in pure BCC Cr. If there are data one can have interaction parameters like `MQ&FE(BCC,Cr,FE)`. Each mobility parameter can depend on T and P.

Other properties that may be described by this model package are lattice parameters, elastic constants, resistivity, etc. see Table 1.

2.4 Components

The term "component" is not used for the modeling, only for controlling the system externally. The term component is reserved for an irreducible set of species that describe the

composition space of the system, sometimes called “system components”. In most cases these components are the elements. The species that can dissolve in a phase are called constituents and can be vacancies, elements, ions or molecules. For most simple models the constituents of the phases are identical to the elements, i.e. to the components.

The components are important as they determine the number of conditions that can be set in order to calculate the equilibrium of the system because one can only set as many conditions as one has components and for T and P. Many conditions, like the amounts or chemical potentials, can only be specified for the components, not for an individual species. This does not mean that the calculation of equilibria is limited to conditions on the amounts of the components, only that the number of components determine the number of conditions that must be set in order to calculate the equilibrium.

In GTP the elements are by default the components but the user may change this to any orthogonal set of species. When working with oxides with no degree of freedom for the oxygen content, like in a quasibinary CaO-SiO₂ system, it may be convenient to select the oxides CaO and SiO₂ as components and then leave the content of the third element undetermined by setting an arbitrary (positive) value of its activity. But in such cases it is necessary to know that the models used for the phases does not allow the content of this element to vary independently. If a gas phase is included in the system it may be more convenient to set the gas as fixed with zero amount, although that may increase the calculation time.

2.5 Fraction sets

In some cases phases with LRO modeled with several sublattices have some properties that are independent of the configuration, only of the composition. In such a case it can be convenient to split the total Gibbs energy in two separate Gibbs energy functions, this is called a “partitioned” or “splitted” model. It was originally used for phases with order/disorder transformations like A1/L1₂ or A2/B2 where the phase can be completely disordered. All parameters to describe the disordered state were collected in a phase that had no sublattices for ordering (it could have a sublattice for interstitials). This made it also easier to include ordering in multicomponent databases where many subsystems have A1 (fcc) or A2 (bcc) phases without ordering.

The Gibbs energy of a partitioned phase used to be written in Thermo-Calc:

$$G_M = G_M^{\text{dis}}(x) + (G_M^{\text{ord}}(y) - G_M^{\text{ord}}(y = x)) \quad (30)$$

$$\Delta G_M^{\text{ord}} = G_M^{\text{4SL}}(y) - G_M^{\text{4SL}}(y = x) \quad (31)$$

The parameters for the ordered phase, as disordered, was included in the disordered phase.

There is no reason to do this with the OC software as the disordered parameters are included in the same phase as the ordered. The Gibbs energy for a partitioned phase is simply:

$$G_M = G_M^{\text{dis}}(x) + G_M^{\text{4SL}}(y) \quad (32)$$

where the parameters in the disordered part only has parameters that describe the disordered contribution, independent of the parameters for the ordering.

A variant of this model have been applied also to intermetallic phases that never disorder completely like σ , μ , Laves phase and similar. The equation is then slightly different as there is no need to describe the completely disordered state

$$G_M = \left(G_M^{\text{dis}}(x) + T^{\text{cfg}} S_M^{\text{dis}}(x) \right) + G_M^{\text{ord}}(y) \quad (33)$$

By adding $T^{\text{cfg}} S_M^{\text{dis}}(x)$ the configurational entropy is only calculated for the ordered part.

This second variant of the disordered model is very interesting because many intermetallic phases like the σ phase have a very large number of end members in multicomponent systems and they must all be given a reasonable value, although in databases today most of them are just equal to the sum of the reference energies of the endmembers, i.e. they can be set to zero. With the increased use of first principles calculations for ordered endmembers, only those which will decrease the Gibbs energy need to be included.

In the OC software we have taken this into account from the beginning and due to the parameters in the disordered part it is not necessary to enter all endmembers of the ordered part, only those which is known to be stable or close to be stable. *Ab initio* data can be used for this when experimental data are missing. The disordered part is entered as an extra “fraction set” of the ordered phase and one simply specifies how many sublattice that should be added together to for the disordered part. For an fcc phase with 5 sublattices, 4 for ordering and one interstitial, one simply gives 4, assuming the 5th sublattice is the interstitial one. For a σ phase modeled with 3 sublattices that all disorder together, one gives 3.

For a ternary σ phase with 5 sublattices one have more than 250 endmembers which all must be calculated. With the new software this may be reduced to less than 10 which will result in a significant increase in calculation speed compared to the case when all endmembers have a value. The maintenance of the database will also be simplified as fewer endmembers are stored.

The new software will create a fraction set record that gives the way to add fractions (the coefficients b_{ij} are stored there as they are also needed to calculate the contribution to the partial derivatives). A link to this fraction set record is stored in the phase_varres record linked from the phase record. When setting the constitution of the phase, i.e. the site fractions, the disordered fractions will be automatically calculated and saved in another phase_varres record linked from the ordered fraction set record. In this way it will thus be straightforward to calculate the ordered and disordered part with the same subroutine, one just have to organize the calculations of each part separately and then add all values, including first and second derivatives, together in the end.

2.6 Composition sets and miscibility gaps

Composition sets is the way OC handles the case when a phase with the same structure can appear simultaneously with two or more compositions. One case this is necessary is when a phase split up into two or more miscibility gaps like the bcc phase in the Cr-Mo system. Less obvious cases is when a phase has an order/disorder transition like the fcc phase in Au-Cu forms L1₂ and L1₀ structures. But also in that case the ordered and disordered phases are described with the same thermodynamic data. Finally this technique is also used to describe the cubic and hexagonal carbides where the metallic atoms are the same as fcc and hcp lattices but interstitial atoms like C and N can occupy most of the available interstitial sites.

Outside the OC package, for example in the TQ package, one can obtain information about the number of composition sets for a phase using the function “nocs(iph)” for each phase “iph”. Inside the OC package the phase number and composition set number are usually given separately. A phase must have at least one composition set and the maximum number is 9. The composition sets larger than 1 is normally suffixed after the phase name separated by a # like BCC#2. If omitted composition set 1 is assumed.

If there are several composition sets of a phase there is a separate phase_varres record for each composition set. Normally these have identical set of constituents but it is possible to suspend constituents in each composition set separately. At the time when a fraction set is added to a phase this must not have any suspended constituents.

Fraction set records have no array or free list, they exist only inside the phase_varres records. (This created some extra headache when implementing them as a simple assignement of a record to another did not create any new record in Fortran08).

2.7 State variables

State variables are, as the name indicate, only dependent on the state of the system, not how this state was reached. In principle they have a well-defined value only at equilibrium but in the modeling they can be extended also outside this range.

The state variables recognized by GTP and that can be used are defined in Table 2. Many of them can have several indices and/or normalizing quantities. In the box below there are some examples of using these in conditions and otherwise.

N=1	means the system has one mole of atoms
N(H)=1	means the system has 1 mole of component H
N(GAS,H)=1	means the gas phase has 1 mole of component H
NP(LIQUID)=1	means the liquid phase has 1 mole of components
X(H)=0.3	means the mole fraction of component H in the system is 0.3
X(GAS,H)=0.3	means the gas phase has a mole fraction of 0.3 of H
B=1000	means the system has a mass of 1000 grams
W%(CR)=18	means the system has 18 mass percent of component CR
W(LIQUID,C)=0.01	means the liquid has 0.01 mass fraction of carbon
H	is the total enthalpy of the system
H(LIQUID)	is the enthalpy of the current amount of liquid (can be zero)
HM(LIQUID)	is the enthalpy of liquid per mole components
HV(LIQUID)	is the enthalpy of liquid per m ³
HW(LIQUID)	is the enthalpy of liquid per gram
HF(LIQUID)	is the enthalpy of one formula unit of liquid

3 Fortran data structures

A number of arrays must be dimensioned, this is done by defining some constants (called PARAMETER in Fortran) and then using these to dimension arrays with fixed size. It may be possible to set these limits in a more flexible way by allocating the arrays in a special subroutine for initiation.

As I am learning Fortran08 by this programming project I discover new possibilities now and again. It seems possible to allow the initialization subroutine to do this dimensioning which means one may tailor the dimensioning of the arrays depending on the kind of calculation one will make.

3.1 User defined data types

The presentation of the data structures defined here follows the way they are declared in the Fortran source code. That may sometimes not be entirely logical but it makes it simpler to update the documentation.

In Fortran08 the programmer can define specific data types and several such are used to store the model information. A data type can contain many different kinds of variables like characters and numerical information. One can also have links between instances (records) of the same and different kinds of datatypes.

This kind of method to organize data is available in all modern computing languages and has been severely missed in Fortran. In Thermo-Calc it was handled by storing data in the self-designed workspace.

Some data types are straightforward to implement like the element. Thus one defines a

Table 2: A very preliminary table with the state variables and their internal representation. Some model parameter properties are also included. The "z" used in some symbols like Sz means the optional normalizing symbol M, W, V or F.

Symbol	Id	Index	Normalizing	Meaning	
		1	2	suffix	
Intensive properties					
T	1	-	-	-	Temperature
P	2	-	-	-	Pressure
MU	3	component	-/phase	-	Chemical potential
AC	4	component	-/phase	-	Activity
LNAC	5	component	-/phase	-	LN(activity)
Extensive properties					
U	10	-/phase#set	-	-	Internal energy for system
UM	11	-/phase#set	-	M	Internal energy per mole
UW	12	-/phase#set	-	W	Internal energy per mass
UV	13	-/phase#set	-	V	Internal energy per m ³
UF	14	phase#set	-	F	Internal energy per formula unit
Sz	2z	-/phase#set	-	-	entropy
Vz	3z	-/phase#set	-	-	volume
Hz	4z	-/phase#set	-	-	enthalpy
Az	5z	-/phase#set	-	-	Helmholtz energy
Gz	6z	-/phase#set	-	-	Gibbs energy
NPz	7z	phase#set	-	-	Moles of phase
BPz	8z	phase#set	-	-	Mass of phase
Qz	9z	phase#set	-	-	Stability of phase
DGz	10z	phase#set	-	-	Driving force of phase
Nz	11z	-/phase#set/comp	-/comp	-	Moles of component
X	111	phase#set/comp	-/comp	0	Mole fraction
X%	111	phase#set/comp	-/comp	100	Mole per cent
Bz	12z	-/phase#set/comp	-/comp	-	Mass of component
W	122	phase#set/comp	-/comp	0	Mass fraction
W%	122	phase#set/comp	-/comp	100	Mass per cent
Y	130	phase#set	const#subl	-	Constituent fraction
Some model parameter identifiers					
TC	-	phase#set	-	-	Curie temperature
BMAG	-	phase#set	-	-	Aver. Bohr magneton number
MQ&X	-	phase#set	constituent	-	Mobility of X
THET	-	phase#set	-	-	Debye temperature

data type which has all the necessary information for an element and then allocates an array to hold as many elements that is necessary.

Other datatypes are more involved, like the record to hold information about an endmember parameter of a phase. This record must refer to the constituents that makes up the endmember and also a link to a list of properties that can be calculated for this endmember. Finally it must be possible to link the endmember record in a list for a specific phase and it must be possible to have a link to interaction records based on this endmember. Before one can figure out how to store endmember data it is thus necessary to have a data structure containing many different kinds of data. One must also consider which data that may be different in different parallel processes dealing with the phase and which data that are “static”. Many of these records may also be of different size for different phases and for example an endmember record for a phase with 3 sublattices need 3 times as many places for constituent indices than a phase with a single lattice. This can be handled with the dynamic allocation feature in Fortran08.

The TYPE definition feature in Fortran08 may be a little less flexible than in other programming languages but it also avoids some of the strange features of these like the “this” operator needed at strange places in some cases. There is no “new” operator either, records are usually declared as single entities or as arrays with fixed (although allocatable) size. But one can use pointers to allocate new records which can be found only by these pointers.

Some of the structures are used to define arrays, like the element, species and phases. Some are created dynamically and can only be accessed by pointers, like endmembers and property records. Some are used locally inside subroutines and declared globally only if used in several.

3.2 Bits of information

Many records contain information that is of YES/NO or ON/OFF type and they are often stored as bits in a status word. Some of these are set automatically and there are subroutines and commands to change others. They are summarized below for all records.

The use of each bit is explained in the text from the source code.

```
!-Bits in global status word (GS) in globaldata record
! level of user: beginner, occational, advanced; NOGLOB: no global gridmin calc
! NOMERGE: no merge of gridmin result,
! NODATA: not any data,
! NOPHASE: no phase in system,
! NOACS: no automatic creation of composition set for any phase
! NOREMCS: do not remove any redundant unstable composition sets
! NOSAVE: data changed after last save command
! VERBOSE: maximum of listing
! SETVERB: permanent setting of verbose
```



```

! SILENT: as little output as possible
! NOAFTEREQ: no manipulations of results after equilibrium calculation
! XGRID: extra dense grid for all phases
! NOPAR: do not run in parallel
! NOSMGLOB do not test global equilibrium at node points
! NOTELCOMP the elements are not the components
! TGRID use grid minimizer to test if global after calculating equilibrium
! OGRID use old grid generator
! NORECALC do not recalculate equilibria even if global test after fails
! OLDMAP use old map algorithm
! NOAUTOSP do not generate automatic start points for mapping
! NO LONGER: HICKEL set if Hickel extrapolation check for solids
! >>>> some of these should be moved to the gtp_equilibrium_data record
integer, parameter :: &
    GSBEG=0,      GSOC=1,      GSADV=2,      GSNOGLOB=3, &
    GSNOMERGE=4,  GSNODATA=5,  GSNOPHASE=6,  GSNOACS=7, &
    GSNOREMCS=8,  GSNOSAVE=9,  GSVERBOSE=10, GSSETVERB=11,&
    GSSILENT=12,  GSNOAFTEREQ=13, GSXGRID=14,  GSNOPAR=15, &
    GSNOSMGLOB=16, GSNOTELCOMP=17, GSTGRID=18,  GSOGRID=19, &
    GSNORECALC=20, GSOLDMAP=21,  GSNOAUTOSP=22
!-----
!-Bits in element record
integer, parameter :: &
    ELSUS=0
!-----
!-Bits in species record
! SUS    Suspended,
! IMSUS  implicitly suspended (when element suspended)
! EL     species is element,
! VA     species is the vacancy
! ION    species have charge,
! SYS    species is (system) component
! UQAC   species used in uniquac model (2 extra reals for area and volume)
integer, parameter :: &
    SPSUS=0, SPIMSUS=1, SPEL=2, SPVA=3, &
    SPION=4, SPSYS=5,  SPUQC=6

```

3.2.1 Phase record bits

The phase record has many bits specifying various things. Additionally the composition set record has other bits, most important if the composition set is ENTERED, meaning it can be stable, FIX, meaning it is a condition that it is stable, SUSPENDED, meaning it must not be stable, and DORMANT meaning it must not be stable but the driving force is calculated.

A user can change these bits but he/she cannot set the CSSTABLE bit which is only set if the phase is stable after a full equilibrium calculation.

```

!-Bits in phase record
! HID phase is hidden (not implemented)
! IMHID phase is implicitly hidden (not implemented)
! ID phase is ideal, substitutional and no interaction
! NOCV phase has no concentration variation (I am not sure it is set)
! HASP phase has at least one parameter entered
! FORD phase has 4 sublattice FCC ordering with parameter permutations
! BORD phase has 4 sublattice BCC ordering with parameter permutations
! SORD phase has TCP type ordering (like for sigma)
! MFS phase has a disordered fraction set
! GAS this is the gas phase (first in phase list)
! LIQ phase is liquid (can be several but listed first after gas)
! IONLIQ phase has ionic liquid model (I2SL)
! AQ1 phase has aqueous model (not implemented)
! DILCE phase has dilute configurational entropy (not implemented)
! QCE phase has quasichemical SRO configurational entropy (not implemented)
! CVMCE phase has some CVM ordering entropy (not implemented)
! EXCB phase need explicit charge balance (has ions)
! XGRID use extra dense grid for this phase
! FACTCE phase has FACT quasichemical SRO model (not implemented)
! NOCS not allowed to create composition sets for this phase
! HELM parameters are for a Helmholtz energy model (not implemented),
! PHNODGDY2 phase has model with no analytical 2nd derivatives
! ELMA phase has elastic model A (not implemented)
! PHSUBO ordering model with ordered part subtracted (is it used??)
! FHV phase has Flory-Huggins model for polymers
! MULTI may be used with care
! BMAV Xion magnetic model with average Bohr magneton number
! UNIQUAC The UNIQUAC fluid model
integer, parameter :: &
  PHHID=0,      PHIMHID=1,  PHID=2,      PHNOCV=3, &      ! 1 2 4 8 : 0/F
  PHHASP=4,     PHFORD=5,   PHBORD=6,    PHSORD=7, &      !
  PHMFS=8,      PHGAS=9,    PHLIQ=10,   PHIONLIQ=11, &    !
  PHAQ1=12,     PHDILCE=13, PHQCE=14,   PHCVMCE=15,&    !
  PHEXCB=16,    PHXGRID=17, PHFACTCE=18, PHNOCS=19,&    !
  PHHELM=20,    PHNODGDY2=21, PHELMA=22, PHSUBO=23,&    !
  PHFHV=24,     PHMULTI=25, PHBMAV=26,  PHUNIQUAC=27   !
!
!-----
!-Bits in constituent fraction (phase_varres) record STATUS2
! CSDFS is set if record is for disordred fraction set, then one must use
!   sublattices from fraction_set record
! CSDLNK: a disordred fraction set in this phase_varres record

```

```

! CSDUM2 and CSDUM3 not used
! CSCONSUS set if one or more constituents suspended (status array constat
!   specify constituent status)
! CSORDER: set if fractions are ordered (only used for BCC/FCC ordering
!   with a disordered fraction set).
! CSABLE: set if phase is stable after an equilibrium calculation ?? needed
! CSAUTO set if composition set created during calculations
! CSDEFCON set if there is a default constitution
! CSTEMPAR set if created by grid minimizer and can be suspended afterwards
!   when running parallel
! CSDEL set if record is not used but has been and then deleted (by gridmin)
! CSADDG means there are terms to be added to G
  integer, parameter :: &
    CSDFS=0,   CSDLNK=1,   CSDUM2=2,   CSDUM3=3, &
    CSCONSUS=4, CSORDER=5, CSABLE=6,   CSAUTO=7, &
    CSDEFCON=8, CSTEMPAR=9, CSDEL=10,   CSADDG=11

```

3.2.2 Some more bits

In particular the `gtp_equilibrium_data` record there are some bits that specifies how the equilibrium calculation can be done, for example if the software can use global grid minimization to find start values.

For the parameter property record there are also bits specifying of the property may depend on T and P or if it has a constituent or component specifier (like the mobility).

```

!-Bits in constat array for each constituent
! For each constituent:
! SUS constituent is suspended (not implemented)
! IMSUS is implicitly suspended,
! VA is vacancy
! QCBOND the constituent is a binary quasichemical cluster
  integer, parameter :: &
    CONSUS=0,   CONIMSUS=1,   CONVA=2,   CONQCBOND=3
!-----
!-Bits in state variable functions (svflista)
! SVFVAL symbol evaluated only explicitly (mode=1 in call) all dot derivatives
! SVFEXT symbol value taken from equilibrium %eqnoval
! SVCONST symbol is a constant (can be changed with AMEND)
! SVFTPF symbol is a TP function, current value returned
! SVFDOT symbol is a DOT function, partial derivative like cp=h.t
  integer, parameter :: &
    SVFVAL=0,   SVFEXT=1,   SVCONST=2,   SVFTPF=3,&
    SVFDOT=4
!-----

```

```

!-Bits in gtp_equilibrium_data record
! EQNOTHREAD set if equilibrium must be calculated before threading
! (in assessment) for example if a symbol must be evaluated in this
! equilibrium before used in another like H(T)-H298
! EQNOGLOB set if no global minimization
! EQNOEQCAL set if no successful equilibrium calculation made
! EQINCON set if current conditions inconsistent with last calculation
! EQFAIL set if last calculation failed
! EQNOACS set if no automatic composition sets ?? not used !! see GSNOACS
! EQGRIDTEST set if grid minimizer should be used after equilibrium
! EQGRIDCAL set if last calculation was using only gridminimizer
  integer, parameter :: &
    EQNOTHREAD=0, EQNOGLOB=1, EQNOEQCAL=2, EQINCON=3, &
    EQFAIL=4, EQNOACS=5, EQGRIDTEST=6, EQGRIDCAL=7
!-----
!-Bits in parameter property type record (gtp_propid)
! no T or P dependence (constant)
! only P dependence
! only T dependence
! there is an element suffix (like mobility),
! there is a constituent suffix
! Property has no addition (used when entering and listing data)
  integer, parameter :: &
    IDNOTP=0, IDONLYP=1, IDONLYT=2, IDELSUFFIX=3, IDCONSUFFIX=4,&
    IDNOADD=5
!-----
!- Bits in condition status word (some set in onther ways??)
! singlevar means T=, x(el)= etc, singlevalue means value is a number
! phase means the condition is a fix phase
  integer, parameter :: &
    ACTIVE=0, SINGLEVAR=1, SINGLEVALUE=2, PHASE=3
!-----
!- Bits in assessment head record status
! ahcoef set means coefficients are entered
  integer, parameter :: &
    AHCOEF=0
!
!-----
!- Bits in addition record status word gtp_phase_add
! havepar set if the phase has parameters for this addition
! if not set the addition is not listed
! permol set if addition should be muliplied with number of atoms
  integer, parameter :: &
    ADDHAVEPAR=0, ADDPERMOL=1
!
! >>> Bits for symbols and TP functions missing ???

```

3.2.3 Phase status revision

A phase status can vary, the default is ENTERED which means that the phase can be stable or not depending on the conditions set by the user. The status values are defined as symbols according to the list below:

```
! some constants, phase status
integer, parameter :: PHHIDDEN=-4
integer, parameter :: PHSUS=-3
integer, parameter :: PHDORM=-2
integer, parameter :: PHENTUNST=-1
integer, parameter :: PHENTERED=0
integer, parameter :: PHENTSTAB=1
integer, parameter :: PHFIXED=2
character (len=12), dimension(-4:2), parameter :: phstate=&
    (/ 'HIDDEN      ', 'SUSPENDEED  ', 'DORMANT      ', 'ENTERED UNST', &
       'ENTERED     ', 'ENTERED STBL', 'FIXED        ' /)
```

The phase status are also available as character strings in the array phstate, defined above, for use in listings.

The HIDDEN status is questionable and not implemented. A phase with the HIDDEN status should not be included in the normal list of phases but stored in a separate list. If the status of a phase is changed to or from being HIDDEN the system should be re-initiated. The possibility to implement HIDDEN is kept as the current database interface does not allow selection of phases when reading from a database.

3.3 Dimensioning

Most arrays are dimensioned in the GTP package using constants. If one need to change them it is usually only in one place. Most of these constants are used in the init_gtp subroutine but also when temporary arrays are needed inside some other subroutines.

```
! Parameters defining the size of arrays etc.
! max elements, species, phases, sublattices, constituents (ideal phase)
! integer, parameter :: maxel=100,maxsp=1000,maxph=600,maxsubl=10,maxconst=1000
! NOTE increasing maxph to 600 and maxtpf to 80*maxph made the equilibrium
! record very big and created problems storing equilibria at STEP/MAP!!!
integer, parameter :: maxel=100,maxsp=1000,maxph=400,maxsubl=10,maxconst=1000
! maximum number of constituents in non-ideal phase
integer, parameter :: maxcons2=100
! maximum number of elements in a species
integer, parameter :: maxspel=10
! maximum number of references
```

```

integer, private, parameter :: maxrefs=1000
! maximum number of equilibria
integer, private, parameter :: maxeq=900
! some dp values, default precision of Y and default minimum value of Y
! zero and one set in tpfun
double precision, private, parameter :: YPRECD=1.0D-6, YMIND=1.0D-30
! dimension for push/pop in calcg, max composition dependent interaction
integer, private, parameter :: maxpp=1000, maxinter=3
! max number of TP symbols, TOO BIG VALUE MAKES SAVE AT STEP/MAP DIFFICULT
integer, private, parameter :: maxtpf=20*maxph
! integer, private, parameter :: maxtpf=80*maxph
! max number of properties (G, TC, BMAG MQ%(...) etc)
integer, private, parameter :: maxprop=50
! max number of state variable functions
integer, private, parameter :: maxsvfun=500
! version number of GTP (not OC)
character*8, parameter :: gtpversion='GTP-3.21'
! THIS MUST BE CHANGED WHENEVER THE UNFORMATTED FILE FORMAT CHANGES!!!
character*8, parameter :: savefile='OCF-3.20'

```

3.3.1 Predefined and user defined additions

The additions provided or added by programmers must be uniquely identified. This is an attempt to do that with those already available. The model parameter identifiers are explained in section 2.3.3 and listed in Table 1.

```

! The number of additions to the Gibbs energy of a phase is increasing
! This is a way to try to organize them. Each addtion has a unique
! number identifying it when created, listed or calculated. These
! numbers are defined here
integer, public, parameter :: INDENMAGNETIC=1
integer, public, parameter :: XIONGMAGNETIC=2
integer, public, parameter :: DEBYECP=3
integer, public, parameter :: EINSTEINCP=4
integer, public, parameter :: TWOSTATEMODEL1=5
integer, public, parameter :: ELASTICMODEL1=6
integer, public, parameter :: VOLMOD1=7
integer, public, parameter :: UNUSED_CRYSTALBREAKDOWNMOD=8
integer, public, parameter :: SECONDEINSTEIN=9
integer, public, parameter :: SCHOTTKYANOMALITY=10
integer, public, parameter :: DIFFCOEFS=11
! name of additions:
character(len=24) , public, dimension(12), parameter :: additionname=&
    ['Inden-Hillert magn model', 'Inden-Xiong magn model ', &
    'Debye CP model', 'Einstein Cp model ', &

```

```

      'Liquid 2-state model      ', 'Elastic model A          ', &
      'Volume model A          ', 'Unused CBT model        ', &
      'Smooth CP step          ', 'Schottky Anomaly       ', &
      'Diffusion coefficients   ', '                        ' ]
!      123456789.123456789.1234  123456789.123456789.1234
! Note that additions often use extra parameters like Curie or Debye
! temperatures defined by model parameter identifiers stored in gtp_propid

```

3.4 The global error code

The error code is the only member of the gtp_parerr record. In parallel execution each thread must have its own value and it is thus declared threadprivate.

```

      TYPE gtp_parerr
! This record contains the global error code. In parallel processing each
! parallel processes has its own error code copied to this if nonzero
! it should be replaced by gtperr for separate errors in treads
      INTEGER :: bmperr
      END TYPE gtp_parerr
      TYPE(gtp_parerr) :: gx
! needed to have error code as private in threads
!$OMP threadprivate(gx)

```

4 Data structures for the TP-fun routines

The TP functions are used to enter model parameters and other functions that depend on T and P. They have a limited syntax as the software must be able to calculate their first and second derivatives with respect to T and P.

The expressions are stored only once but the calculated values of the functions are stored separately for each equilibria as each equilibria can have different values of T and P.

4.1 Structure to store expressions of TP functions

In this record the coefficients and powers of T and P and possible unary functions used is stored. There is a simple parser that can read a simple TP function similar to what TC has. My idea has been to extend this a bit by allowing parenthesis in a more flexible way but that is not yet implemented. A TP function can refer to another TP function using the link array. This means a recursive evaluation of TP functions must be implemented.

This information must not be changed during parallel processing. During assessments one may calculate each experimental equilibria in parallel and as the coefficients of TP

functions are varied by the assessment procedure one may think that this array must also be separate for each parallel process. But changes of coefficients are not done during the parallel execution so there is no problem.

```
integer, parameter :: tpfun_expression_version=1
  TYPE tpfun_expression
! Coefficients, T and P powers, unary functions and links to other functions
    integer noofcoeffs,nextfrex
    double precision, dimension(:), pointer :: coeffs
! each coefficient kan have powers of T and P/V and links to other TPFUNS
! and be multiplied with a following LOG or EXP term.
! wpow USED FOR MULTIPLYING WITH ANOTHER FUNCTION!!
    integer, dimension(:), pointer :: tpow
    integer, dimension(:), pointer :: ppow
    integer, dimension(:), pointer :: wpow
    integer, dimension(:), pointer :: plevel
    integer, dimension(:), pointer :: link
  END TYPE tpfun_expression
! These records are allocated when needed, not stored in arrays
```

4.2 Bits used in TP function status

These bits are used for different purposes

```
! BITS in TPFUN
! TPCONST      set if a constant value
! TPOPTCON     set if optimizing value
! TPNOTENT     set if referenced but not entered (when reading TDB files)
! TPVALUE      set if evaluated only explicitly (keeping its value)
integer, parameter :: &
    TPCONST=0,    TPOPTCON=1,    TPNOTENT=2,    TPVALUE=3
```

4.3 Function root record type

In this record a name of the function is stored and the number of temperature ranges. For each range a low temperature limit and a link to a tpfun_expression record is stored. There is also a high temperature limit. As the number of ranges can vary the arrays are allocatable (declared as pointers). At the same time as a root record is reserved to store a TP function one also reserves a tpfun_parres record and in this the last calculated result of the function is stored. Saving the last calculated values speeds up calculation because the same function is often used in many parameters and needed many times for the same values of T and P. But in parallel processing the values of T and P can be different in each processor and thus the results must be separate in each.


```

integer, parameter :: tpfun_root_version=1
TYPE tpfun_root
! Root of a TP function including name with links to coefficients and codes
! and results. Note that during calculations which can be parallelized
! the results can be different for each parallel process
character*(lenfnsym) symbol
! Why are limits declared as pointers?? They cannot be properly deallocated
! limits are the low temperature limit for each range
! funlinks links to expression records for each range
! each range can have its own function, status indicate if T and P or T and V
! forcenewcalc force new calculation when optimizing variable changed
integer noofranges,nextfree,status,forcenewcalc
double precision, dimension(:), pointer :: limits
TYPE(tpfun_expression), dimension(:), pointer :: funlinks
double precision hightlimit
END TYPE tpfun_root
! These records are stored in arrays as the actual function is global but each
! equilibrium has its own result array (tpfun_parres) depending on the local
! values of T and P/V. The same index is used in the global and local arrays.
! allocated in init_gtp
TYPE(tpfun_root), private, dimension(:), pointer :: tpfuns

```

4.4 Structure for calculated results of TP functions

This record contains the last values of T and P used for calculation the TP function and the results including first and second derivatives of T and P. These are ordered as F, F.T, F.P, F.T.T, F.T.P and F.P.P. The reason to have this as a separate array is that in parallel processing the values of T and P may be different in each process and thus the results also, whereas the values of the coefficients are the same. The size of all the function records are allocated dynamically by the tpfun.init subroutine.

```

integer, parameter :: tpfun_parres_version=1
TYPE tpfun_parres
! Contains TP results, 6 double for results and 2 doubles for T and P
! values used to calculate the results
! Note that during calculations which can be parallelized the
! results can be different for each tread
integer forcenewcalc
double precision, dimension(2) :: tpused
double precision, dimension(6) :: results
END TYPE tpfun_parres
! This array is local to the gtp_equilibrium_data record
! index is the same as the function

```

5 Basic thermodynamic data structures

Below the data types in GTP will be explained and in some cases also how they are declared as arrays or inside other data types. Many of the declared variables are “private” which means they cannot be changed directly from outside the module. For such data that the user will manipulate there are subroutines or functions provided.

5.1 Global data

Information that is valid for the whole system is stored here. Note that T and P may be different in different parallel processes and if there is a prescribed value that is stored in a condition record. Maybe this is not really needed.

```
TYPE gtp_global_data
! status should contain bits how advanced the user is and other defaults
! it also contain bits if new data can be entered (if more than one equilib)
! sysparam are variables for different things
! sysparam(1) unused
! sysparam(2) number of equilibria between each check of spinodal at STEP/MAP??
! sysparam(3-10) unused ...
! sysreal(1) is the minimum T for EET check (equi-entropy T, Hickel)
!           if zero no EET c
      integer status
      integer :: sysparam(10)=0
      character name*24
      double precision rgas,rgasuser,pnorm
      double precision :: sysreal(5)=zero
END TYPE gtp_global_data
TYPE(gtp_global_data) :: globaldata
```

5.2 Version identification of the data structure

As the data structure is always open to changes and as such changes may affect other parts of the software most of the TYPE definitions have a “version” value. Whenever a change is made in a TYPE definition this version value should be incremented. The version number can be tested in other parts of the software, in particular when writing the data structure on a file and at a later time reading it back from a file. If the version number is not the same it may not be possible to read the file or the software must have provisions for reading records with different version numbers.

5.3 The ELEMENT data type

Elements are the building blocks of other data. They have a symbol, mass, reference state and some other values defined.

The elements in a system are stored in an array ELLISTA of these records in the order they are entered. Another integer array ELEMENTS have the elements in alphabetical order. These arrays are allocated in the subroutine init_gtp.

There are two predefined elements, the electron with symbol “/-” and the vacancy with symbol “Va”. They have the indices -1 and 0. All real elements have numbers from 1 and higher.

The OC software is case insensitive, UPPER and lower case letters are treated identically. Thus Va, va and VA is the same. Elements with a single letter symbol must be followed by a space or a non-alphabetic character, for example a stoichiometric factor, to separate it from a following element.

The advantage with ELLISTA is that the element is stored at an index which never change, it does not change when other elements are entered which would change the alphabetical order. Note that there is a cross index stored in ellista so it is possible to know the alphabetical order of the element.

It also means there are 3 ways to specify an element, its symbol, its alphabetical index (its index in ELEMENTS) and its index in ELLISTA. The ELLISTA index is most important as it will never change whereas the ELEMENTS index may change when new elements are entered which may change the alphabetical order. The link to an element from a species are always the ELLISTA index.

In the species record the stoichiometric formula is stored by giving the index of the element in ELLISTA. The elements array is useful to make alphabetically ordered listing of data. Elements are normally never deleted but may be suspended i.e. hidden from application programs (this cannot be allowed during parallel processing).

```
! this constant must be incremented whenever a change is made in gtp_element
INTEGER, parameter :: gtp_element_version=1
TYPE gtp_element
! data for each element: symbol, name, reference state, mass, h298-h0, s298
    character :: symbol*2,name*12,ref_state*24
    double precision :: mass,h298_h0,s298
! splink: index of corresponding species in array splink
! Status bits are stored in the integer status
! alphaindex: the alphabetical order of this elements
! refstatesymbol: indicates H0 (1), H298 (0, default) or G (2) for endmembers
    integer :: splink,status,alphaindex,refstatesymbol
END TYPE gtp_element
! allocated in init_gtp
TYPE(gtp_element), private, allocatable :: ellista(:)
```

```
INTEGER, private, allocatable :: ELEMENTS(:)
```

5.4 Species

Here some of the useful features of Fortran08 data structuring appears and can be explained. The species have a name, mass and charge as fixed attributes. It also has a status word and a link back to the array where the species are arranged in alphabetical order. It has an integer giving the number of elements in the species (must be larger than zero). Finally there are two arrays which have no size specified in the declaration, these must be allocated when the species is entered. The reason to have arrays that can be allocated is that some species has just one element, other may have two, three or more. The dimension of these arrays are stored in `noofel` (It can actually also be obtained by the Fortran08 build-in function `SIZE`). The indices stored in `ellinks` are the location, i.e. index to `ellista`. This does not change even if the alphabetical order of elements is changed.

For saving and reading a stored datastructure from a file it is necessary to have the number of elements as a stored integer because this is needed to allocate space for these arrays before reading the content of these arrays from the file. The `SIZE` function is useless in that case.

One can declare arrays that should be allocated dynamically either as `ALLOCATABLE` or `POINTER`. Note that allocating `POINTER` variables is a likely source of memory leaks.

The species record is an interesting example of an array of structures that in itself contains allocatable arrays that can vary in size depending on the data stored there. This feature will be utilized extensively when storing data for phases.

For the `UNIQUAC` model an allocatable array, `spextra`, with double precision variables has been added in version 5 of `OC`. This allows storing the area and volume parameter associated with a component in the `UNIQUAC` model.

It is also a very interesting feature to use indices to other arrays as links rather than pointers which is necessary in most other languages with data structures. There are pointers also in Fortran08 but pointers have a bad habit of being confusing and complicated to use, one is never sure if one has the address of a pointer or the content (which is also a memory address).

The integer array “species” has the species in alphabetical order by providing the correct index to the `SPLISTA`.

```
! this constant must be incremented whenever a change is made in gtp_species
INTEGER, parameter :: gtp_species_version=2
TYPE gtp_species
! data for each species: symbol, mass, charge, extra, status
! mass is in principle redundant as calculated from element mass
    character :: symbol*24
    double precision :: mass, charge
```

```

! alphaindex: the alphabetical order of this species
! noofel: number of elements
! nextra: number of extra properties (size of spextra)
      integer :: noofel,status,alphaindex
! Use an integer array ellinks to indicate the elements in the species
! The corresponding stoichiometry is in the array stoichiometry
      integer, dimension(:), allocatable :: ellinks
      double precision, dimension(:), allocatable :: stoichiometry
! Can be used for extra species properties as in UNIQUAC models (area, volume)
      double precision, dimension(:), allocatable :: spextra
END TYPE gtp_species
! allocated in init_gtp
TYPE(gtp_species), private, allocatable :: splista(:)
INTEGER, private, allocatable :: SPECIES(:)

```

5.5 Components

It will be possible to define different components for each equilibrium. A component must be a species. Initially the species identical to the elements are the components. The components are part of the "gtp_equilibrium_data" record and after a calculation the chemical potentials are stored in these records.

```

! this constant must be incremented whenever a change is made in gtp_component
INTEGER, parameter :: gtp_component_version=1
TYPE gtp_components
! The components are simply an array of indices to species records
! the components must be "orthogonal". There is always a set of "systems
! components" that by default is the elements.
! Later one may implement that the user can define a different "user set"
! and maybe also specific sets for each phase.
! The reference state is set as a phase and value of T and P.
! The name of the phase and its link and the link to the constituent is stored
! the endmember array is for the reference phase to calculate GREF
! The last calculated values of the chemical potentials (for user defined
! and default reference states) should be stored here.
! molat is the number of moles of components in the defined reference state
      integer :: splink,phlink,status
      character*16 :: refstate
      integer, dimension(:), allocatable :: endmember
      double precision, dimension(2) :: tpref
      double precision, dimension(2) :: chempot
      double precision mass,molat
END TYPE gtp_components
! allocated in gtp_equilibrium_data

```

5.6 Phase datatypes

The data stored for a phase is very complex. All data are accessed from a phase “root” record (except when save/read from a file). The phase has one or more phase_varres records which is stored in the gtp_equilibrium_data record. Each equilibrium has such a record and in these records the last calculated results are kept. The user can create several equilibria with different sets of conditions as thus the results will be different in each equilibrium. This is a way to handle parallel processing but it is also useful to store experimental data in assessments.

In the phase_varres record all fraction variables (and some additional information) that can be different for each parallel process is also stored. If the phase has two or more composition sets each of these has its own phase_varres record. The phase_varres records are an array inside the gtp_equilibrium_data record and each phase store the links to its composition sets by indices to this array. This is necessary as one can have several gtp_equilibrium_data records and one must have the same number of composition sets for a phase in all equilibria. Whenever new composition sets are created in parallel computing it must be done for all threads at the same time.

The model parameter data is the same for all equilibria, it is part of the “static” data structure. The root record of a phase has data and links to all necessary information listed below:

- The endmember record list. There can be two endmember lists, one for each fraction types.
- The phase_varres record representing a composition set. Each phase can have up to 9 composition sets. The phase_varres records are declared as an array in the gtp_equilibrium_data record because each equilibrium must have a unique set of fractions and results. In this record there are also some status bits that may be different in different parallel processes like if the phase (composition set) is FIXED/ENTERED/etc.

A phase may exist simultaneously with different constituent fractions like in a miscibility gap or when a phase can order. To identify a composition set one can use the number symbol “#” followed by a digit. When the composition set is created one can also add a pre- and suffix.

In the phase_varres record there are also arrays to store all calculated G and its first and second derivatives (also derivatives with respect to fractions). Calculated values of other properties like TC, V, MQ etc (and their derivatives) are also stored if there are parameters for these properties.

For the ionic liquid model the number of sites depend on the constitution and thus there are also provisions to have the number of sites on each sublattice in this record.

- The addition list. This is a pointer to an addition record (there can be several linked sequentially). The only current addition record is used to specify the magnetic mag-

netic contribution and for other additions new subroutines must be written to enter, list and calculate the additions. See section 5.6.9.

These records may in turn point to other records, as already mentioned the endmember record has links to another endmember records and to an interaction records and both of these records has a link to a list of property records where there are links to TP functions. Some of these links are indexes to arrays of these records but for example the parameters stored in endmember, interactions and property records are created dynamically and can only be accessed by pointers. The property records have links to TP functions but these links are indices to the array of TP function structure. The reason to have an array for these is that there are separate arrays with the calculated results of the TP functions in the `gtp-equilibrium_data` record as these must be local to each equilibrium, whereas the function expressions are the same for all equilibria.

5.6.1 Permutations of constituents

A complication for endmembers and interaction records is to handle permutations of constituents when one has ordering. It is clumsy to store identical permutation of constituents (like A:A:A:B, A:A:B:A, A:B:A:A, B:A:A:A for a 4 sublattice FCC ordering) in separate endmembers but one should have a single endmember for all permutations. It is necessary that the software can handle such permutations for multicomponent systems, otherwise the databases must contain several 1000 permutations of identical values of the same parameter. A 4 component FCC ordered system with 4 sublattices have totally 256 endmembers but only 35 unique ones.

The permutations create complications also for the interaction records which depend on the order of constituents in the endmember. An interaction records for an interaction between A and B belonging to an end member A:A:A:A (which has no permutations) but there are 4 permutations of the interaction record (B can be in any of the 4 sublattices). So each combination must be considered.

The second order interaction A,B:A,C:A:A this has 3 additional permutations compared to the first order as the interaction with C can be placed in any of the 3 remaining sublattices. But the second order interaction A,B:A,B:A:A has a more complicated permutations. If the first interaction with B is in the first sublattice, the second interaction with B can be placed in any of the remaining 3 but if the first level interaction in B is in the second sublattice the second level interaction can only be in the third or forth. In Table 3 some permutations are shown.

Reciprocal parameters like $L(\text{fcc}, A, B: A, B: A: A)$ are important to approximate the short range order contributions to the Gibbs energy.

Table 3: Possible permutations of some FCC endmembers and interaction parameters. In the table the component arrays within parenthesis represent a permutation. There are no permutations of a parameter $G(\text{FCC}, \text{A:A:A:A})$ but 4 permutations of the parameter $L(\text{FCC}, \text{A,B:A:A:A})$ and 6 permutations of $L(\text{FCC}, \text{A,B:A,B:A:A})$. The 2nd level permutations are linked from the corresponding 1st level in order to increase the speed of calculation.

endmember	1st level interaction	2nd level interaction
A: A: A: A	A,B: A: A: A	A,B: A,B: A: A (A,B: A: A,B: A) (A,B: A: A: A,B) (A: A,B: A,B: A) (A: A,B: A: A,B) (A: A: A,B: A) (A: A: A: A,B) no permutation
A: A: A: B	A,B: A: A: B	A,B: A,B: A: B (A,B: A: A,B: B) (A: A,B: A,B: B) no permutation
(A: A: B: A)	(A: A: A,B: B) (A,B: A: B: A)	(A,B: A,B: B: A) (A,B: A: B: A,B) (A: A,B: B: A) (A: A: B: A,B) no permutation
etc.		

In order to enter and modify parameters that are stored as permutation it is essential to have the constituents in a strict order. Of course the user is allowed to enter the components for each sublattice in any order but the program will rearrange them according to there rules, assuming a system like

(A B C D)(A B C D)(A B C D)(A B C D):

- The alphabetically first component will be in sublattice 1 and the other ordered in increasing order. For option B where the first two and the following two are different the order applies to each pair of sublattices. Thus a parameter entered as $G(\text{BCC4}, \text{D:B:C:A})$ will be sorted as $G(\text{A:C:B:A})$.
- For interactions the alphabetically first constituent will be included in the endmember. Interactions will be moved to the first possible sublattice. For reciprocal parameters with interactions in two sublattices and when the endmember component is the same the alphabetically first interacting component will come in a sublattice before the other. For example $L(\text{BCC}, \text{B:C:A}, \text{C:A,B})$ will be sorted as $L(\text{BCC}, \text{A,B:A}, \text{C:B:C})$

Without such strict ordering of constituents it will not be possible to find a parameter in the data structure.

Table 4: Som more permutations of some endmembers and interaction records for BCC. In this case the endmember, 1st and 2nd interactions have the same number of permutations.

endmember	1st level interaction	2nd level interaction
A: B: A: C	A,B: B: A: C	A,B: B: A,C: C
(B: A: A: C)	(B: A,B: A: C)	(B: A,B: A,C: C)
(A: B: C: A)	(A,B: B: C: A)	(A,B: B: C: A,C)
(A: C: A: B)	(A: C: A,B: B)	(A,C: C: A,B: B)
(C: A: A: B)	(C: A: A,B: B)	(C: A,B: A,C: B)
(C: A: B: A)	(C: A: B: A,B)	(C: A,C: B : A,B)
etc.		

The permutations means some complications when entering parameters as links to the fractions of all permutations are created at that stage. It also means that all constituents must be ordered, always in alphabetical order, so it is possible to find a parameter if its value should be changed. When calculating with the permutations the data structure created when entering the parameter must contain information on the number of permutations inherited from the endmember up to the second level interaction. No interactions higher than the 2nd order are allowed for a phase with permutations.

Since the first release of OC most of the permutations for the tetrahedron 4 sublattice fcc (and hcp) ordering has been implemented. The 4 sublattice bcc permutations are more complex but are now also implemented. All endmember permutations and all first level level have been implemented but only a limited set of 2nd order interactions, i.e. all for binary and ternary system. See also sections 8.6.9 and 8.6.10

An ordered phase will in many cases have a disordered fraction set for the disordered state and there will be parameters which depend on these fractions. There can be endmembers and interaction parameters for each fraction set. The parameters for each fraction set will be evaluated separately and added together at the end, taking into account the chain rule for the derivatives.

5.6.2 Property types for different fraction sets

An endmember or interaction can have several property records, normally one for the Gibbs energy but also for Curie temperature, mobilities etc. But there can be endmembers and interaction records without property records just because some higher interaction record have a property record.

For mobilities one can use a special type of property that is specific to a constituent like MQ&FE. In a property record one must have an adjustable array for function links, for example when there are several RK terms, see 2.3.2. These links can be allocated dynamically

and if there are new RK terms added or deleted this array can be extended or decreased dynamically. For each property record one can store a reference to the origin of the data.

I am not sure exactly how one should in listings identify parameters belonging to different fraction sets. In a SIGMA phase with 3 sublattices one will have endmember parameters for some like G(SIGMA,FE:CR:FE), which are multiplied with the site fractions of these constituents, $y_{1,Fe}y_{2,Cr}y_{3,Fe}$. But we will also have a fraction set, which in this case, use the mole fractions of Fe and Cr,

$$x_{Fe} = (10y_{1,Fe} + 4y_{2,Fe} + 16y_{3,Fe})/30$$

An endmember parameter for this fraction set is currently specified as GD(SIGMA,FE) and an interaction as GD(SIGMA,CR,FE). If in the future one is interested to extend to several levels of fraction sets one should maybe use an integer specification like G#2(SIGMA,FE) etc. but that seems unlikely for the next 10 years at least. In any cases a property without specification will belong to the primary site fraction set which is the major one, all other fractions are calculated from the fractions in the primary fraction set.

For FCC ordering with interstitials the second fraction set will not be mole fractions but the sum of the site fractions of the substitutional sublattices

$$z_{Fe} = 0.25 \sum_{s=1,4} y_{s,Fe}$$

5.6.3 Selected_element_reference

With Thermo-Calc it has been a little awkward that one may not read the normal reference phase for an element from the database. For example the gas phase may not be included for calculations with nitrogen present but one would like to have partial pressures or activities of nitrogen referenced to the gas.

An attempt to handle this has been made by entering by default a phase called “SELECT_ELEMENT_REFERENCE”. This phase is always hidden and cannot be included in any calculation. Its location and index is 0 (zero) so it does not show up in the number of phases. But the user can list the data of this phase and also amend them. The idea is that a database will automatically enter the appropriate data for each element in this phase and in calculations this phase will be the default reference state for activities or enthalpies (at current temperature). One must have a phase and not just a TP function to handle elements with magnetic transitions. This reference phase is in fact a different phase for each element, one must be able to use different magnetic functions for different elements. All details for this are not worked out and it may not even be a very good idea.

5.6.4 The endmember record

As already described the phase record is a root of two lists of endmembers, one for each of the two possible fraction sets. The endmember records are allocated by pointers when needed and contains several links to other records:

- one link to the next endmember record,
- one link to an interaction record, the root of a binary tree,
- one link to a list of property records and
- links to one constituent in each sublattice

All the links except those to the constituents may be empty. The links to the constituents are not real links but an integer index of the array of constituents stored in the phase record (described later). The endmembers are always arranged in increasing value of these indices, in the order of the sublattices. The endmember properties are multiplied with the fractions of these constituents when a calculation is performed.

When the ordering options has been implemented, see 5.6.1, an endmember may have several permutations of the constituent indices. That is the reason to have two-dimensional arrays for the constituent indices.

One may have endmembers that does not depend on the constituent in a specific sublattice and this is represented by the fraction index for this sublattice is negative, usually -99. This is called a “wildcard” and represented by an asterix “*” when entering or listing the parameter, see section 2.3.4.

```
! this constant must be incremented whenever a change is made in gtp_endmember
INTEGER, parameter :: gtp_endmember_version=1
TYPE gtp_endmember
! end member parameter record, note ordered phases can have
! several permutations of fraction pointers like for B2: (Al:Fe) and (Fe:Al).
! There are links (i.e. indices) to next end member and to the interactio tree
! and to a list of property record
! The phase link is needed for SAVE/READ as one cannot know the number of
! sublattices otherwise. One could just store nsl but a link back to the
! phase record might be useful in other cases.
! noofpermut: number of permutations (for ordered phases: (Al:Fe) and (Fe:Al))
! phaselink: index of phase record
! antalem: sequential order of creation, useful to keep track of structure
! propointer: link to properties for this endmember
! nextem: link to next endmember
! intponter: root of interaction tree of parameters
! fraclinks: indices of fractions to be multiplied with the parameter
    integer :: noofpermut,phaselink,antalem
    TYPE(gtp_property), pointer :: propointer
    TYPE(gtp_endmember), pointer :: nextem
    TYPE(gtp_interaction), pointer :: intpointer
! there is at least one fraclinks per sublattice
! the second index of fraclinks is the permutation (normally only one)
```

```

! the first indec of fraclinks points to a fraction for each sublattice.
! The fractions are numbered sequentially independent of sublattices, a
! sigma phase with (FE:CR,MO:CR,FE,MO) has 6 fractions (incl one for FE in
! first sublattice) and the end member (FE:MO:CR) has the fraclinks 1,3,4
! This means these values can be used as index to the array with fractions.
! The actual species can be found via the sublattice record
!   integer, dimension(:,:), pointer :: fraclinks
!   integer, dimension(:,:), allocatable :: fraclinks
END TYPE gtp_endmember
! dynamically allocated when entering a parameter

```

5.6.5 The interaction record

The interaction records forms a binary tree starting from an endmember record. It thus has two pointers to other interaction records, one which is the *next* on the same interaction level, excluding the interacting constituent in the current record, and one to a *higher* level, including the current interacting constituent. In this way any level of interaction parameter can be defined. However, only the first two levels (binary and ternary) can be composition dependent, see 2.3.2.

There is also a link to a list of property records. This link can be empty if there is a link to a higher order interaction.

The interaction record specifies one additional constituent in a sublattice. The remaining constituents are given by the endmember record and any lower interaction records with a higher link to this one. Again one can have permutations of this interacting constituent in the different sublattices, see section 5.6.1.

The interaction records are always linked from the first possible endmember, i.e. that with the lowest set of constituent indices in all sublattices. If there are several levels of interactions the lowest interaction has the lowest constituent index. There is no ordering of interacting constituents on the same level.

```

! this constant must be incremented when a change is made in gtp_interaction
INTEGER, parameter :: gtp_interaction_version=1
TYPE gtp_interaction
! this record constitutes the parameter tree. There are links to NEXT
! interaction on the same level (i.e. replace current fraction) and
! to HIGHER interactions (i.e. includes current fraction)
! There can be several permutations of the interactions (both sublattice
! and fraction permuted, like interaction in B2 (Al:Al,Fe) and (Al,Fe:Al))
! The number of permutations of interactions can be the same, more or fewer
! compared to the lower order parameter (endmember or other interaction).
! The necessary information is stored in noofip. It is not easy to keep
! track of permutations during calculations, the smart way to store the last
! permutation calculated is in this record ... but that will not work for

```

```

! parallel calculations as this record is static ...
! status: may be useful eventually
! antalint: sequential number of interaction record, to follow the structure
! order: for permutations one must have a sequential number in each node
! propointer: link to properties for this parameter
! nextlink: link to interaction on same level (replace interaction)
! highlink: link to interaction on higher level (include this interaction)
! sublattice: (array of) sublattices with interaction fraction
! fraclink: (array of) index of fraction to be multiplied with this parameter
! noofip: (array of) number of permutations, see above.
      integer status,antalint,order
      TYPE(gtp_property), pointer :: propointer
      TYPE(gtp_interaction), pointer :: nextlink,highlink
      integer, dimension(:), allocatable :: sublattice,fraclink,noofip
END TYPE gtp_interaction
! allocated dynamically and linked from endmember records and other
! interaction records (in a binary tree)

```

5.6.6 The property record

The endmembers and interaction records is the start of a list of property records. A property list can be empty if an endmember or interaction record is needed to specify some higher interaction.

There is a property index, see 2.3.8 and a *nextpr* link to another property record. One can have a link to a reference for the parameter (published paper or similar), see 5.6.7.

The actual value of the property is stored as an index to a TP function. These functions are stored in array and cannot be deleted (but they can be changed) so the index is a stable link. The calculated values of a property is stored in another array, local to the equilibrium record, see 6.2.5, as one may have different values in each thread in parallel processing or when running assessments, as these use different equilibrium records.

For binary and ternary interactions one can have a degree larger than zero which means there are several TP functions linked to this property. See 2.3.2.

```

! this constant must be incremented when a change is made in gtp_property
INTEGER, parameter :: gtp_property_version=1
TYPE gtp_property
! This is the property record. The end member and interaction records
! have pointer to this. Several different properties can be linked
! from a parameter record like G, TC, BMAGN, VA, MQ etc.
! Some properties are connected to a constituent (or component?) like the
! mobility and also the Bohr mangneton number.
! Allocated as linked from endmembers and interaction records
! reference: can be used to indicate the source of the data

```

```

! refix: can be used to indicate the source of the data
! nextpr: link to next property record
! extra: TOOP and KOHLER can be implemented inside the property record
! proptype: type of property, 1 is G, other see parameter property
! degree: if parameter has Redlich-Kister or similar degrees (powers)
! degreeLink: indices of TP functions for different degrees (0-9)
! protect: can be used to prevent listing of the parameter
! antalprop: probably redundant (from the time of arrays of property records)
      character*16 reference
      TYPE(gtp_property), pointer :: nextpr
      integer proptype,degree,extra,protect,refix,antalprop
      integer, dimension(:), allocatable :: degreeLink
END TYPE gtp_property
! property records, linked from endmember and interaction records, allocated
! when needed. Each property like G, TC, has a property record linking
! a TPFUN record (by index to tpfun_parres)

```

5.6.7 Bibliographic references

It is important to document the source of the data and each property can have a unique bibliographic reference stored in this record. It would typically be a published paper or some internal written documentation.

```

! this constant must be incremented when a change is made in gtp_biblioref
! old name gtp_datareference
      INTEGER, parameter :: gtp_biblioref_version=1
      TYPE gtp_biblioref
! store data references
! reference: can be used for search of reference
! refspect: free text
      character*16 reference
!      character*64, dimension(:), allocatable :: refspect
! this is Fortran 2003/2008 standard, not available in GNU 4.8
!      character(len=:), allocatable :: nyrefspect
! Use wpack routines!!!
      integer, dimension(:), allocatable :: wprefspect
END TYPE gtp_biblioref
! allocated in init_gtp
      TYPE(gtp_biblioref), private, allocatable :: bibrefs(:)

```

5.6.8 Parameter property identification

The property list has an index and the meaning of this index is given the gtp_propid record. A few of these are predefined but a programmer can add more, see section 2.3.3. For any

added property software must also be written to handle such properties during calculations. The implemented properties can be listed, see 8.7.26 and the values of these properties can be listed similarly like for state variables, see 8.7.27.

```
! this constant must be incremented when a change is made in gtp_propid
INTEGER, parameter :: gtp_propid_version=1
TYPE gtp_propid
! this identifies different properties that can depend on composition
! Property 1 is the Gibbs energy and the others are usually used in
! some function to contribute to the Gibbs energy like TC or BMAGN
! But one can also have properties used for other things like mobilities
! with additional especification like MQ&FE
! symbol: property identifier like G for Gibbs energy
! note: short description for listings
! prop_elsymb: additional for element dependent properties like mobilities
      character symbol*4,note*28,prop_elsymb*2
! Each property has a unique value of idprop. Status can state if a property
! has a constituent specifier or if it can depend on T or P
      integer status
! this can be a constituent specification for Bohr mangetons or mobilities
! such specification is stored in the property record, not here
!      integer prop_spec,listid
! >>> added "listid" as a conection to the "state variable" listing here.
! This replaces TC, BMAG, MQ etc included as "state variables" in order to
! list their values. In this way all propids become available
end TYPE gtp_propid
! the value TYPTY stored in property records is "idprop" or
! if IDELSUFFIX set then 100*"idprop"+ellista index of element
! if IDCONSUFFIX set then 100*"idprop"+constituent index
! When the parameter is read the suffix symbol is translated to the
! current element or constituent index
TYPE(gtp_propid), dimension(:), private, allocatable :: propid
```

5.6.9 Additions to the Gibbs energy

Different contributions to the Gibbs energy can be specified in the software using the data structure below. Such additions normally depend on a number of composition dependent properties like the Curie temperature, the Bohr magneton number, the Debye temperature etc. These can be added as properties and entered as endmember and interaction parameters with a unique index and symbol see 5.6.8.

When implementing a new addition subroutines must be written for entering, listing and calculating the property, including analytical first and second derivatives with respect to T , P and the constitution.

One may also add properties that does not contribute to the Gibbs energy but depend on the constitution of the phase, like mobilities, viscosities, resistivity etc. Such properties are calculated together with the Gibbs energy and their values can be extracted by appropriate subroutines.

As mentioned in section 1.4 most addition model calculate a Gibbs energy contribution per atom whereas the Gibbs energy model for the phase need the value per mole formula unit. There is a bit ADDPERMOL for this in the status word for additions defined in section 3.2.2.

```
! this constant must be incremented when a change is made in gtp_phase_add
  INTEGER, parameter :: gtp_phase_add_version=1
  TYPE gtp_phase_add
! record for additions to the Gibbs energy for a phase like magnetism
! addrecno: ?
! aff: antiferromagnetic factor (Inden model)
! status: BIT 1 set if there are parameters
! need_property: depend on these properties (like Curie T)
! explink: function to calculate with the properties it need
! nextadd: link to another addition
    integer type,addrecno,aff,status
    integer, dimension(:), allocatable :: need_property
    TYPE(tpfun_expression), dimension(:), pointer :: explink
    TYPE(gtp_phase_add), pointer :: nextadd
    type(gtp_elastic_modela), pointer :: elastica
    type(gtp_diffusion_model), pointer :: diffcoefs
! calculated contribution to G, G.T, G.P, G.T.T, G.T.P and G.P.P
    double precision, dimension(6) :: propval
  END TYPE gtp_phase_add
! allocated when needed and linked from phase record
```

5.6.10 The elastic model

This is a tentative record to model elastic contributions to the Gibbs energy.

```
! addition record to calculate the elastic energy contribution
! declared as allocatable in gtp_phase_add
! this constant must be incremented when a change is made in gtp_elastic_modela
  INTEGER, parameter :: gtp_elastic_modela_version=1
  TYPE gtp_elastic_modela
! lattice parameters (configuration) in 3 dimensions
    double precision, dimension(3,3) :: latticepar
! epsilon in Voigt notation
    double precision, dimension(6) :: epsa
! elastic constant matrix in Voigt notation
    double precision, dimension(6,6) :: cmat
```



```

! calculated elastic energy addition (with derivative to T and P?)
    double precision, dimension(6) :: eeadd
! maybe more ...
end TYPE gtp_elastic_modela

```

5.6.11 The diffusion coefficients

This is a record to handle mobilities and diffusion coefficients. So far it has not been used in any application.

```

! addition record to calculate diffusion coefficients
! declared as allocatable in gtp_phase_add
! this constant must be incremented when a change is made in gtp_elastic_modela
    INTEGER, parameter :: gtp_diffusion_model_version=1
    TYPE gtp_diffusion_model
! status bit 0 set means no calculation of this record
! dilute, simple or magnetic
        integer difftypemodel,status
! alpha values for magnetic diffusion (for interstitials in constituent order)
        double precision, allocatable, dimension(:) :: alpha
! indices of dependent constituent in each sublattices
        integer, allocatable, dimension(:) :: depcon
! indices of constituents with zero volume
        integer, allocatable, dimension(:) :: zvcon
! calculated diffusion matrix
        double precision, allocatable, dimension(:, :) :: dcoef
! Maybe we need one for each composition set?? at least to save the matrix
        type(gtp_diffusion_model), pointer :: nextcompset
! maybe more ...
end TYPE gtp_diffusion_model

```

5.6.12 Data structures to transform TDB to DAT files

With OC it is possible to convert a TDB file to a SOLGASMIX dat file. These are data structures used for that.

```

    TYPE gtp_tpfun_as_coeff
! this is a TPFUN converted to coefficients without any references to other
! functions. Each function can have several T ranges and coefficients for T**n
        double precision, dimension(:), allocatable :: tbreaks
        double precision, dimension(:, :), allocatable :: coefs
        integer, dimension(:, :), allocatable :: tpows
! this is used only during conversion
!     type(gtp_tpfun_as_coeff), pointer :: nextcrec

```

```

    end type gtp_tpfun_as_coeff
!
    INTEGER, parameter :: gtp_tpfun2dat_version=1
    TYPE gtp_tpfun2dat
! this is a temporary storage of TP functions converted to arrays of
! coefficients. Allocated as an array when necessary and the index in
! this array is the same index as for the TPfun
        integer nranges
!         type(gtp_tpfun_as_coeff) :: tpfuncoef
        type(gtp_tpfun_as_coeff) :: cfun
    end type gtp_tpfun2dat

```

5.6.13 The phase and composition set indices

In many cases one must specify both a phase and a composition set and this is done by separate indices in most subroutine calls. As this is not very convenient an attempt to introduce a single index for both phase and composition sets has been made by the “phase tuple” type. This is a simple record with five integers, ixphase is the index of the phase in alphabetical order, compset the composition set index, phaseix is the index of the phase in the phlista array, lokvares is the index of the phase_varres record which is useful to retrieve data. If there is a higher composition set for the same phase then nextcs is the phase tuple index to the phase, otherwise nextcs is zero. Ixphase is probably redundant. There is an array with the phasetuple records called PHASETUPLE which is updated whenever a composition set is created or deleted.

```

! this constant must be incremented when a change is made in gtp_phasetuple
    INTEGER, parameter :: gtp_phasetuple_version=1
    TYPE gtp_phasetuple
! for handling a single array with phases and composition sets
! ixphase is phase index, compset is composition set index
! ADDED also index in phlista (lokph) and phase_varres (lokvares) and
! nextcs which is nonzero if there is a higher composition set of the phase
! A tuple index always refer to the same phase+compset. New tuples with
! the same phase and other compsets are added at the end.
! BUT if a compset>1 is deleted tuples with higher index will be shifted down!
        integer lokph,compset,ixphase,lokvares,nextcs
! >>>>>>>>>>>> old      integer phaseix,compset,ixphase,lokvares,nextcs
    end TYPE gtp_phasetuple

```

5.6.14 The phase record

We have finally reached the phase record itself. As this refers to many of the structures above it is declared in the source code after all of them and to simplify the updating of this

documentation we follow the order of the declarations in the source code. It is complicated enough already ...

The phase record had originally two parts but these have been merged to a single record. The link to the composition set data (the phase_varres record) has also recently been changed so the link to all of them are stored in the phase record as an array of indices to the array of phase_varres records in the equilibrium record. The phase_varres record contains all *dynamic* data that change during iterations and has also all calculated results. The endmember and interaction records contain the *static* data that does not change (except the calculated values of all TP functions⁹ but this is also stored the equilibrium record in the tp_res array.

When a phase is created a link to a record in the phase_varres array, i.e. an integer index, is stored in the array linktocs(1). If additional composition sets are created they are stored sequentially in the same array. A phase cannot have more than 9 composition sets.

```
! a smart way to have an array of pointers used in gtp_phase
TYPE endmemrecarray
    type(gtp_endmember), pointer :: p1
end TYPE endmemrecarray
!-----
! this constant must be incremented when a change is made in gtp_phase
INTEGER, parameter :: gtp_phase_version=1
TYPE gtp_phaserecord
! this is the record for phase model data. It points to many other records.
! Phases are stored in order of creation in phlista(i) and can be found
! in alphabetical order through the array phases(i)
! sublista is now removed and all data included in phlista
! sublattice and constituent data (they should be merged)
! The constituent link is the index to the splista(i), same function
! as LOKSP in iws. Species in alphabetical order is in species(i)
! One can allocate a dynamic array for the constituent list, done
! by subroutine create_constitlist.
! Note that the phase has a dynamic status word status2 in gtp_phase_varres
! which can be different in different parallel calculations.
! This status word has the FIX/ENT/SUS/DORM status bits for example
! name: phase name, note composition sets can have pre and suffixes
! model: free text
! phletter: G for gas, L for liquid
! alphaindex: the alphabetical order of the phase (gas and liquids first)
    character name*24,models*72,phletter*1
    integer status1,alphaindex
! noofcs: number of composition sets,
! nooffs: number of fraction sets (replaces partitioned phases in TC)
    integer noofcs,nooffs
! additions: link to addition record list
! ordered: link to endmember record list
```

```

! disordered: link to endmember list for disordered fractions (if any)
  TYPE(gtp_phase_add), pointer :: additions
  TYPE(gtp_endmember), pointer :: ordered,disordered
! To allow parallel processing of endmembers, store a pointer to each here
  integer noemr,ndemr
  TYPE(endmemrecarray), dimension(:), allocatable :: oendmemarr,dendmemarr
! noofsubl: number of sublattices
! tnooffr: total number of fractions (constituents)
! linktocs: array with indices to phase_varres records
! nooffr: array with number of constituents in each sublattice
! Note that sites are stored in phase_varres as they may vary with the
! constitution for ionic liquid)
  integer noofsubl,tnooffr
  integer, dimension(9) :: linktocs
  integer, dimension(:), allocatable :: nooffr
! number of sites in phase_varres record as it can vary with composition
! constitlist: indices of species that are constituents (in all sublattices)
  integer, dimension(:), allocatable :: constitlist
! used in ionic liquid:
! i2slx(1) is index of Va, i2slx(2) is index of last anion (both can be zero)
  integer, dimension(2) :: i2slx
! allocated in init_gtp.
  END TYPE gtp_phaserecord
! NOTE phase with index 0 is the reference phase for the elements
! allocated in init_gtp
  TYPE(gtp_phaserecord), private, allocatable :: phlista(:)
  INTEGER, private, allocatable :: PHASES(:)

```

5.7 State variables and the state variable record

Conditions and results are obtained as values of state variables. There are many of these like G, H, T, x (< component >) etc. They are stored in the software as a record of the type below where *istv* is an integer giving the basic type and indices can give additional specification. Some properties like chemical potentials can have a reference state and one can also define a unit like Kelvin or calories. At present this is used only to specify if a composition variable is a fraction or a percent.

```

! this constant must be incremented when a change is made in gtp_state_variable
  INTEGER, parameter :: gtp_state_variable_version=1
  TYPE gtp_state_variable
! this is to specify a formal or real argument to a function of state variables
! statev/istv: state variable index >=9 is extensive
! phref/iref: if a specified reference state (for chemical potential
! unit/iunit: 100 for percent, no other defined at present

```

```

! argtyp together with the next 4 integers represent the indices(4), only 0-4
! argtyp=0: no indices (T or P)
! argtyp=1: component
! argtyp=2: phase and compset
! argtyp=3: phase and compset and component
! argtyp=4: phase and compset and constituent
! ?? what is norm ?? normalizing like M in HM ?
      integer statevarid,norm,unit,phref,argtyp
! these integers represent the previous indices(4)
      integer phase,compset,component,constituent
! a state variable can be part of an expression with coefficients
! the coefficient can be stored here. Default value is unity.
! In many cases it is ignored
      double precision coeff
! NOTE this is also used to store a condition of a fix phase
! In such a case statev is negative and the absolute value of statev
! is the phase index. The phase and compset indices are also stored in
! "phase" and "compset" ??
! This is a temporary storage of the old state variable identifier
      integer oldstv
      end TYPE gtp_state_variable
! used for state variables/properties in various subroutines

```

6 Use of the thermodynamic data structures

This section describes how the basic thermodynamic data structures are used in various applications like calculating the Gibbs energy, handling errors, step for property diagrams, mapping of phase diagram, assessments etc.

6.1 Error handling

There is a global error code defined as part of the defined in the TYPE gtp-parerr. This has just one integer variable called bmperr and there is one variable of this type called gx which is declared THREADPRIVATE for use in parallel processing.

The error code is gx%bmperr in the whole OC system. Whenever there is an error gxraising the error and usually the subroutine is exited directly. The error code should be tested after each subroutine call and maybe the calling routine can handle the error.

There are error messages defined for the errors generated by GTP.

6.2 Calculations

So far we have described how to handle the data for the phase. In order to calculate an equilibrium many values, like the constituent fractions, amounts of phase, temperature etc may change. As mentioned several times we must also be able to handle several separate equilibria, either as threads in parallel computing or as experimental data in assessments. Each of these separate datasets are stored in a `gtp_equilibrium_data` record.

6.2.1 Conditions

We must also handle conditions set by the user. Typically a condition is a state variable equal to a value like $T=1273$ or $w(c)=0.01$. At present simple assignments of state variables are the main types of conditions allowed but for phase compositions it is possible to use expressions like $x(\text{liq},\text{fe})-x(\text{bcc},\text{fe})=0$ to find a congruent and to set a state variables equal to a symbolic value like $w(\text{liq},b)=\text{whatever}$.

```
! this constant must be incremented when a change is made in gtp_condition
! NOTE on unformatted SAVE files the conditions are written as texts
  INTEGER, parameter :: gtp_condition_version=1
  TYPE gtp_condition
! these records form a circular list linked from gtp_equilibrium_data records
! each record contains a condition to be used for calculation
! it is a state variable equation or a phase to be fixed
! The state variable is stored as an integer with indices
! NOTE: some state variables cannot be used as conditions: Q=18, DG=19, 25, 26
! There can be several terms in a condition (like  $x(\text{liq},c)-x(\text{fcc},c)=0$ )
! nofterms: number of terms in condition expression
! statev: the type of state variable (must be the same in all terms)
!         negative value of statev means phase index for fix phase
! active: zero if condition is active, nonzero for other cases
! unit: is 100 if value in percent, can also be used for temperature unit etc.
! nid: identification sequential number (in order of creation), redundant
! iref: part of the state variable (iref can be comp.set number)
! iunit: ? confused with unit?
! seqz is a sequential index of conditions, used for axis variables
! experimtype: inequality (< 0 or > 0) and/or percentage (-101, 100 or 101)
! symlink: index of symbol for prescribed value (1) and uncertainty (2)
! condcoeff: there is a coefficient and set of indices for each term
! prescribed: the prescribed value
! NOTE: if there is a symlink value that is the prescribed value
! current: the current value (not used?)
! uncertainty: the uncertainty (for experiments)
      integer :: nofterms,statev,active,iunit,nid,iref,seqz,experimenttype
!     TYPE(putfun_node), pointer :: symlink1,symlink2
```

```

! better to let condition symbol be index in svflista array
  integer symlink1, symlink2
  integer, dimension(:,:), allocatable :: indices
  double precision, dimension(:), allocatable :: condcoeff
  double precision prescribed, current, uncertainty
! confusing with record statevar and integer statev
  TYPE(gtp_state_variable), dimension(:), allocatable :: statvar
  TYPE(gtp_condition), pointer :: next, previous
end TYPE gtp_condition
! used inside the gtp_equilibrium_data record and elsewhere

```

6.2.2 State variable functions

In this record the description of a state variable function is stored. The actual expression is stored using the PUTFUN subroutine in the metlib package. The calculated results of a state variable function is stored as a double precision array svfunres in the equilibrium data record. State variable function values are a single value, not 6 as for the TPFuns, as one cannot calculate a derivative with respect to anything.

The “dot” derivative expression available in Thermo-Calc has been implemented for one kind of calculations, $H.T$ meaning the partial derivative of enthalpy with respect to temperature, normally this is the heat capacity. Such a property can only be calculated in connection with an equilibrium calculation or a property diagram.

```

! this constant must be incremented when a change is made in gtp_putfun_list
  INTEGER, parameter :: gtp_putfun_list_version=1
  TYPE gtp_putfun_list
! these are records for state variable functions. The function itself
! is handled by the putfun package.
! linknode: pointer to start node of putfun expression
! narg: number of symbols in the function
! nactarg: number of actual parameter specifications needed in call
!   (like @P, @C and @S
! status: can be used for various things
! status bit SVFVAL=1 means value evaluated only when called with mode=1
! SVCONST bit set if symbol is just a constant value (linknode is zero)
! eqnoval: used to specify the equilibrium the value should be taken from
!   (for handling what is called "variables" in TC, SVFEXT set also)
! SVFTPF set if symbol is a TP function, eqnoval is TPFUN index
! name: name of symbol
  integer narg, nactarg, status, eqnoval
  type(putfun_node), pointer :: linknode
  character name*16
! THIS IS NOT REALLY USED, VALUES ARE STORED IN CEQ%SVFUNRES
  double precision svfv

```

```

! this array has identification of state variable (and other function) symbols
    integer, dimension(:,:), pointer :: formal_arguments
end TYPE gtp_putfun_lista
! this is the global array with state variable functions, "symbols"
TYPE(gtp_putfun_lista), dimension(:), allocatable :: svflista
! NOTE the value of a function is stored locally in each equilibrium record
! in array svfunres.
! The number of entered state variable functions. Used when a new one stored
integer, private :: nsvfun

```

6.2.3 Fraction sets

A phase can have several composition sets, meaning that it can be stable with two or more different compositions. A phase can also have two fraction sets, explained in more detail in 5.6.2.

```

! this constant must be incremented when a change is made in gtp_fraction_set
INTEGER, parameter :: gtp_fraction_set_version=1
TYPE gtp_fraction_set
! info about disordered fractions for some phases like ordered fcc, sigma etc
! latd: the number of sublattices added to first disordred sublattice
! ndd: sublattices for this fraction set,
! tnoofxfr: number of disordered fractions
! tnoofyfr: same for ordered fractions (=same as in phlista).
! varreslink: index of disordered phase_varres,
! totdis: 0 indicates no total disorder (sigma), 1=fcc, bcc or hcp
! id: parameter suffix, D for disordered
! dsites: number of sites in sublattices, disordred fractions stored in
!         another phase_varres record with index varreslink (above)
! splink: pointers to species record for the constituents
! nooffr: the number of fractions in each sublattice
! y2x: the conversion from sublattice constituents to disordered and
! dxidyj: are the the coeff to multiply the y fractions to get the disordered
!         xfra(y2x(i))=xfra(y2x(i))+dxidyj(i)*yfra(i)
! disordered fractions stored in the phase_varres record with index varreslink
! arrays originally declared as pointers now changed to allocatable
    integer latd,ndd,tnoofxfr,tnoofyfr,varreslink,totdis
    character*1 id
    double precision, dimension(:), allocatable :: dsites
    integer, dimension(:), allocatable :: nooffr
    integer, dimension(:), allocatable :: splink
    integer, dimension(:), allocatable :: y2x
    double precision, dimension(:), allocatable :: dxidyj
! formula unit factor needed when calculating G for disordered sigma etc
    double precision fsites

```



```

END TYPE gtp_fraction_set
! these records are declared in the phase_varres record as DISFRA for
! each composition set and linked from the phase_varres record

```

6.2.4 The phase_varres record for composition sets

Each composition set of a phase has a record as described below. It contains all data that can vary during calculations like the constituent fractions, the amount of the phase, etc. It also contains all results from a calculation. An array phase_varres is allocated in the gtp_equilibrium_record for this purpose and in the phase record there are indices to the phase_varres records for its composition sets.

As composition sets can be created and deleted there is a free list maintained using the integer *nextfree*. The first free phase_varres record is given by the global variable *csfree*. This list is maintained in the first equilibrium record, which is pointed to by the global variable *FIRSTEQ*.

```

! this constant must be incremented when a change is made in gtp_phase_varres
! added quasichemical bonds
INTEGER, parameter :: gtp_phase_varres_version=2
TYPE gtp_phase_varres
! Data here must be different in equilibria representing different experiments
! or calculated in parallel or results saved from step or map.
! nextfree: In unused phase_varres record it is the index to next free record
!   The global integer csfree is the index of the first free record
!   The global integer highcs is the highest varres index used
! phlink: is index of phase record for this phase_varres record
! status2: has composition set status bits CSxyz
! phstate: indicate state: fix/stable/entered/unknown/dormant/suspended/hidden
!           2    1    0    -1    -2    -3    -4
! phtupx: phase tuple index
      integer nextfree,phlink,status2,phstate,phtupx
! abnorm(1): moles of components per formula unit of the phase/composition set
! abnorm(2): mass of components per formula unit
! abnorm(3): moles atoms per formula unit (all abnorm set by SET_CONSTITUTION)
! prefix and suffix are added to the name for composition sets 2 and higher
      double precision, dimension(3) :: abnorm
      character*4 prefix,suffix
! constnat: array with status word for each constituent, any can be suspended
! yfr: the site fraction array
! mmyfr: min/max fractions, negative is a mininum
! sites: site ratios (which can vary for ionic liquids)
      integer, dimension(:), allocatable :: constnat
      double precision, dimension(:), allocatable :: yfr
      real, dimension(:), allocatable :: mmyfr
      double precision, dimension(:), allocatable :: sites
! for ionic liquid derivatives of sites wrt fractions (it is the charge),

```

```

! 2nd derivates only when one constituent is vacancy
! 1st sublattice P=\sum_j (-v_j)*y_j + Qy_Va
! 2nd sublattice Q=\sum_i v_i*y_i
! dpqdy is the abs(valency) of the species, set in set_constitution
! for the vacancy it is the same as the number of sites on second subl.
! used in the minimizer and maybe elsewhere
    double precision, dimension(:), allocatable :: dpqdy
    double precision, dimension(:), allocatable :: d2pqdvay
! disfra: a structure describing the disordered fraction set (if any)
! for extra fraction sets, better to go via phase record index above
! this TYPE(gtp_fraction_set) variable is a bit messy. Declaring it in this
! way means the record is stored inside this record.
    type(gtp_fraction_set) :: disfra
! ---
! stored calculated results for each phase (composition set)
! amfu: is amount formula units of the composition set (calculated result)
! netcharge: is net charge of phase
! dgm: driving force
! qcbonds: quasichemical bonds (NOT SAVED ON UNFORMATTED)
    double precision amfu,netcharge,dgm,qcbonds
! qcsro: current value of SRO (for quasichemical model)
    double precision, allocatable, dimension(:) :: qcsro
! Other properties may be that: gval(*,2) is TC, (*,3) is BMAG, see listprop
! nprop: the number of different properties (set in allocate)
! listprop(1): is number of calculated properties
! listprop(2:listprop(1)): identifies the property stored in gval(1,ipy) etc
! 2=TC, 3=BMAG. Properties defined in the gtp_propid record
    integer nprop
    integer, dimension(:), allocatable :: listprop
! gval etc are for all composition dependent properties, gval(*,1) for G
! gval(*,1): is G, G.T, G.P, G.T.T, G.T.P and G.P.P
! dgval(1,j,1): is first derivatives of G wrt fractions j
! dgval(2,j,1): is second derivatives of G wrt fractions j and T
! dgval(3,j,1): is second derivatives of G wrt fractions j and P
! d2gval(ixsym(i,j),1): is second derivatives of G wrt fractions i and j
    double precision, dimension(:,:), allocatable :: gval
    double precision, dimension(:,:,:), allocatable :: dgval
    double precision, dimension(:,:), allocatable :: d2gval
! added for strain/stress, current values of lattice parameters
    double precision, dimension(3,3) :: curlat
! saved values from last equilibrium for dot derivative calculations
    double precision, dimension(:,:), allocatable :: cinvy
    double precision, dimension(:), allocatable :: cxmol
    double precision, dimension(:,:), allocatable :: cdxmol
! terms added to G if bit CSADDG nonzero
    double precision, dimension(:), allocatable :: addg

```

```

! integer containing the iteration when invsaved updated
    integer invsavediter
! arrays to save time in calc_dgdyterms, do not need to be saved on unformatted
    double precision, dimension(:,:), allocatable :: invsaved
    END TYPE gtp_phase_varres
! this record is created inside the gtp_equilibrium_data record

```

6.2.5 The equilibrium record

The equilibrium record has all data that may change dynamically during a calculation. One may have several equilibrium records and during parallel computing each thread must have one. Also during assessments each experimental data is stored in a separate equilibrium record as it has its unique set of conditions.

```

! this must be incremented when a change is made in gtp_equilibrium_data
    INTEGER, parameter :: gtp_equilibrium_data_version=1
    TYPE gtp_equilibrium_data
! this contains all data specific to an equilibrium like conditions,
! status, constitution and calculated values of all phases etc
! Several equilibria may be calculated simultaneously in parallel threads
! so each equilibrium must be independent
! NOTE: the error code must be local to each equilibria!!!!
! During step and map each equilibrium record with results is saved
! values of T and P, conditions etc.
! Values here are normally set by external conditions or calculated from model
! local list of components, phase_varres with amounts and constitution
! lists of element, species, phases and thermodynamic parameters are global
! status: not used yet?
! multiuse: used for various things like direction in start equilibria
! eqno: sequential number assigned when created
! next: index of next free equilibrium record
!      also index of next equilibrium in a list during step/map calculation.
! eqname: name of equilibrium
! comment: a free text, for example reference for experimental data.
! tpval(1) is T, tpval(2) is P, rgas is R, rtn is R*T
! rtn: value of R*T
! weight: weight value for this experiment, default unity
!      integer status,multiuse,eqno,next
!      integer status,multiuse,eqno,nexteq
!      character eqname*24,comment*72
!      double precision tpval(2),rtn
!      double precision :: weight=one
! svfunres: the values of state variable functions valid for this equilibrium
!      double precision, dimension(:), allocatable :: svfunres
! the experiments are used in assessments and stored like conditions

```

```

! lastcondition: link to condition list
! lastexperiment: link to experiment list
    TYPE(gtp_condition), pointer :: lastcondition,lastexperiment
! components and conversion matrix from components to elements
! complist: array with components (species index or location)??
! compstoi: stoichiometric matrix of compoents relative to elements
! invcompstoi: inverted stoichiometric matrix
    TYPE(gtp_components), dimension(:), allocatable :: complist
    double precision, dimension(:,:), allocatable :: compstoi
    double precision, dimension(:,:), allocatable :: invcompstoi
! one record for each phase+composition set that can be calculated
! phase_varres: here all calculated data for the phases are stored
    TYPE(gtp_phase_varres), dimension(:), allocatable :: phase_varres
! index to the tpfun_parres array is the same as in the global array tpres
! eq_tpres: here local calculated values of TP functions are stored
! should be allocatable, not a pointer
    TYPE(tpfun_parres), dimension(:), allocatable :: eq_tpres
! current values of chemical potentials stored in component record but
! duplicated here for easy acces by application software
    double precision, dimension(:), allocatable :: cmuval
! xconv: convergence criteria for constituent fractions and other things
! dgconv(1) is controlling decrease of DGM for unstable phases
! dgconv(2) not used (yet)
    double precision xconv,gdconv(2)
! delta-G value for merging gridpoints in grid minimizer
! smaller value creates problem for test step3.OCM, MC and austenite merged
    double precision :: gmindif=-5.0D-2
! testing merging again 190604/BoS
    double precision :: gmindif=-1.0D-2
! maxiter: maximum number of iterations allowed
    integer maxiter
! This is to store additional things not really invented yet ...
! It may be used in ENTER MANY_EQUIL for things to calculate and list
    character (len=80), dimension(:), allocatable :: eqextra
! this is to save a copy of the last calculated system matrix, needed ??
! to calculate dot derivatives, initiate to zero
    integer :: sysmatdim=0,nfixmu=0,nfixph=0
    integer, allocatable :: fixmu(:)
    integer, allocatable :: fixph(:,:)
    double precision, allocatable :: savesysmat(:,:)
END TYPE gtp_equilibrium_data
! The primary copy of this structures is declared globally as FIRSTEQ here
! Others may be created when needed for storing experimental data or
! for parallel processing. A global array of these are
TYPE(gtp_equilibrium_data), dimension(:), allocatable, target :: eqlista
TYPE(gtp_equilibrium_data), pointer :: firsteq

```

```

! This array of equilibrium records are used for storing results during
! STEP and MAP calculations.
TYPE(gtp_equilibrium_data), dimension(:), allocatable :: eqlines

```

6.3 Records with data shared by several subroutines

The records below have no global variables but are used in some of the calculating subroutines to store complex temporary data, almost like an old-fashioned COMMON area but it is declared inside a subroutine and passed as an argument to the different subroutines.

6.3.1 Parsing data

The data in this record is used parsing the endmember lista and the binary interaction tree.

```

! for each permutation in the binary interaction tree of an endmember one must
! keep track of the permutation and the permutation limit.
! It is not possible to push the value on pystack as one must remember
! them when changing the endmember permutation
! integer, parameter :: permstacklimit=150
! this constant must be incremented when a change is made in gtp_parcalc
INTEGER, parameter :: gtp_parcalc_version=1
TYPE gtp_parcalc
! This record contains temporary data that must be separate in different
! parallel processes when calculating G and derivatives for any phase.
! There is nothing here that need to be saved after the calculation is finished
! global variables used when calculating G and derivaties
! sublattice with interaction, interacting constituent, endmember constituents
! PRIVATE inside this structure not liked by some compilers....
! endcon must have maxsubl dimension as it is used for all phases
integer :: intlat(maxinter),intcon(maxinter),endcon(maxsubl)
! interaction level and number of fraction variables
integer :: intlevel,nofc
! interacting constituents (max 4) for composition dependent interaction
! iq(j) indicate interacting constituents
! for binary RK+Muggianu iq(3)=iq(4)=iq(5)=0
! for ternary Muggianu in same sublattice iq(4)=iq(5)=0
! for reciprocal composition dependent iq(5)=0
! for Toop, Kohler and simular iq(5) non-zero (not implemented)
integer :: iq(5)
! fraction variables in endmember (why +2?) and interaction
double precision :: yfrem(maxsubl+2),yfrint(maxinter)
! local copy of T, P and RT for this equilibrium
double precision :: tpv(2),rgast
! double precision :: ymin=1.0D-30

```

```

    end TYPE gtp_parcalc
! this record is declared locally in subroutine calcg_nocheck

```

6.3.2 Fraction product stack

The product of the constituent fractions and their derivatives must be saved now and again during the parsing. The record below is used for that.

```

! this constant must be incremented when a change is made in gtp_pystack
INTEGER, parameter :: gtp_pystack_version=1
TYPE gtp_pystack
! records created inside the subroutine push/pop_pystack
! data stored during calculations when entering an interaction record
! previous: link to previous record in stack
! ipermutsave: permutation must be saved
! intrecsave: link to interaction record
! pysave: saved value of product of all constituent fractions
! dpysave: saved value of product of all derivatives of constituent fractions
! d2pysave: saved value of product of all 2nd derivatives of constit fractions
    TYPE(gtp_pystack), pointer :: previous
    integer :: pmqsave
    TYPE(gtp_interaction), pointer :: intrecsave
    double precision :: pysave
    double precision, dimension(:), allocatable :: dpysave
    double precision, dimension(:), allocatable :: d2pysave
end TYPE gtp_pystack
! declared inside the calcg_internal subroutine

```

6.4 STEP and MAP results data structures

Some application software which is closely related to the equilibrium calculation, like STEP and MAP have some of their data structures declared here as they would otherwise not be able to access the data.

6.4.1 The node point record

In the gtp_eqnode record an equilibrium representing a node point with several lines meeting is stored. It has links to other node points and to two or more lines of calculated equilibria.

```

INTEGER, parameter :: gtp_eqnode_version=1
TYPE gtp_eqnode
! This record is to arrange calculated equilibria, for example results
! from a STEP or MAP calculation, in an ordered way. The equilibrium records

```

```

! linked from an eqnode record should normally represent one or more lines
! in a diagram but may be used for other purposes.
! ident is to be able to find a specific node
! nodedtype is to specify invariant, middle, end etc.
! status can be used to suppress a line
! color can be used to specify color or linetypes (dotted, thick ... etc)
! exits are the number of lines that should exit from the node
! done are the number of calculated lines currently exiting from the node
      integer ident,nodedtype,status,color,exits,done
! this node can be in a multilayered list of eqnodes
      type(gtp_eqnode), pointer :: top,up,down,next,prev
! nodeq is a pointer to the equilibrium record at the node
      type(gtp_equilibrium_data), pointer :: nodeq
! eqlista are pointers to line of equilibria starting or ending at the node
! The equilibrium records are linked with a pointer inside themselves
      type(gtp_equilibrium_data), dimension(:), pointer :: eqlista
! axis is the independent axis variable for the line, negative means decrement
! noeqs gives the number of equilibria in each eqlista, a negative value
! indicates that the node is an endpoint (each line normally has a
! start point and an end point)
      integer, dimension(:), allocatable :: axis,noeqs
! This is a possibility to specify a status for each equilibria in each line
!      integer, dimension(:,:), allocatable :: eqstatus
end TYPE gtp_eqnode
! can be allocated in a gtp_applicationhead record

```

6.5 Assessment records

Multicomponent thermodynamic databases are created by assessments of many binary and ternary systems based on descriptions of the pure elements. In an assessment experimental and theoretical data are fitted by model parameters for the individual phases. In order to create large databases and extrapolate to multicomponent systems the data and models used for the lower order systems, in particular the pure elements, must be identical.

The assessment of a phase requires that one can include experimental data in the data structure and by using an optimizing software that can vary some of the model parameters. The necessary data structure for this is provided in the GTP package.

The setup of an assessment will include the commands (note the commands listed below may change in later versions of the user interface):

1. ENTER elements, species, phases etc. usually from a macro file.
2. ENTER OPT to enter the coefficients to be optimized. Part of the TPfun package.
3. ENTER PARAMETERS with coefficients to be optimized.

4. ENTER EQUILIBRIA with experimental data (possibly from a file). The experimental data for each equilibrium added with ENTER EXPERIMENT command, see section 8.8.5
5. SET RANGE_EXP_EQUIL to specify the equilibria with experiments.
6. SET VARIABLE_COEFF to specify coefficients to be optimized.
7. OPTIMIZE to make a least square fit.
8. LIST OPTIMIZATION to list the result
9. SAVE TDB to create a TDB file with the results (need editing).
10. and other commands as necessary

6.5.1 The assessment head record

This is the record organizing an assessment. It has all the values related to the optimizing coefficients and has a list of pointers to the equilibria with experimental data. An instance of this is created when starting the software and thus accessible in all packages that use GTP, the arrays declared inside the record can be updated by such packages. There is a possibility to create a linked list of these records to save temporary versions of an assessment. This is not yet implemented and some of the other variables in this record are not yet used.

The status word has just a single bit defined, AHCOEF, set if the optimizing coefficients have been entered by the command ENTER OPTIMIZE_COEFF. The assessment coefficients are TP function constants named A00 to A99. These TP constants can be used entering other TP functions and phase parameters that should be optimized. There are also subroutines to change the values of these variables from an optimizing software. In order to control the optimization the user can also provide a scaling factor, a minimum and a maximum of a coefficient.

The eqlista array contains pointers to the equilibria with experimental data to be used in the assessment. This is set by a command in the user interface, SET RANGE_EXP_EQUIL. The equilibria with experiments are simply entered with an ENTER EQUILIBRIUM command and for each equilibrium the associated experimental data is entered with the command ENTER EXPERIMENT.

```
! a smart way to have an array of pointers used in gtp_assessmenthead
TYPE equilibrium_array
    type(gtp_equilibrium_data), pointer :: p1
end TYPE equilibrium_array
INTEGER, parameter :: gtp_assessment_version=1
TYPE gtp_assessmenthead
! This record should summarize the essential information about assessment data
! using GTP. How it should link to other information is not clear.
! status is status word, AHCOEF is used
! varcoef is the number of variable coefficients
! firstexpeq is the first equilibrium with experimental data
```



```

! lwam is allocated workspace at last call to lmdif1
    integer status,varcoef,firstexpeq,lwam
    character*64 general,special
    type(gtp_assessmenthead), pointer :: nextash,prevash
! This is list of pointers to equilibria to be used in the assessment
! size(eqlista) is the number of equilibria with experimental data
    type(equilibrium_array), dimension(:), allocatable :: eqlista
! These are the coefficients values that are optimized,
! current values, scaling, start values, RSD and optionally min and max
    double precision, dimension(:), allocatable :: coeffvalues
    double precision, dimension(:), allocatable :: coeffscale
    double precision, dimension(:), allocatable :: coeffstart
    double precision, dimension(:), allocatable :: coeffrsd
    double precision, dimension(:), allocatable :: coeffmin
    double precision, dimension(:), allocatable :: coeffmax
! These are the corresponding TP-function constants indices
    integer, dimension(:), allocatable :: coeffindex
! This array indicate currently optimized variables:
! -1=unused, 0=fix, 1=fix with min, 2=fix with max, 3=fix with min and max
! 10=optimized, 11=opt with min, 12=opt with max, 13=opt with min and max
    integer, dimension(:), allocatable :: coeffstate
! Work arrays ...
    double precision, dimension(:), allocatable :: wopt
end TYPE gtp_assessmenthead
! this record should be allocated for assessments when necessary
type(gtp_assessmenthead), allocatable :: ashrecord
! type(gtp_assessmenthead), pointer :: firstash,lastash
! but this is later allocated, to avoid memory loss ashrecord should be used
! and then this pointer should be set to that record
type(gtp_assessmenthead), pointer :: firstash

```

6.6 Other application head record

This is a template for other applications that need access to the internal data of OC.

```

INTEGER, parameter :: gtp_applicationhead_version=1
TYPE gtp_applicationhead
! This record should summarize the essential information about an application
! using GTP. How it should link to other information is not clear.
! The character variables should be used to indicate that.
    integer apptyp,status
    character*64 general,special
! These can be used to define axis and other things
    integer, dimension(:), allocatable :: ivals
    double precision, dimension(:), allocatable :: rvals

```

```

        character*64, dimension(:), allocatable :: cvals
        type(gtp_applicationhead), pointer :: nextapp,prevapp
! The headnode can be the start of a structure of eqnodes with lines
        type(gtp_eqnode) :: headnode
! this is the start of a list of nodes with calculated lines or
! single equilibria that belong to the application.
        type(gtp_eqnode), dimension(:), allocatable :: nodlista
    end TYPE gtp_applicationhead
! this record is allocated when necessary
    type(gtp_applicationhead), pointer :: firstapp,lastapp

```

7 Global variables

The variables below contain some information used in several subroutines. Additionally many of the record types described earlier are declared as private arrays to protect them somewhat.

```

! counters for elements, species and phases initiated to zero
    integer, private :: noofel=0,noofsp=0,noofph=0
! counter for phase tuples (combination of phase+compset)
    integer, private :: nooftuples=0
! counters for property and interaction records, just for fun
    integer, private :: noofprop,noofint,noofem
! free lists in phase_varres records and addition records
    integer, private :: csfree,addresc
! free list of references and equilibria
    integer, private :: reffree,eqfree
! maximum number of properties calculated for a phase
    integer, private :: maxcalcprop=20
! highcs is highest used phase_varres record (for copy equil etc)
    integer, private :: highcs
! Trace for debugging (not used)
    logical, private :: ttrace
! Output for debugging gridmin
    integer, private :: lutbug=0
! minimum constituent fraction
    double precision :: bmpymin
! number of defined property types like TC, BMAG etc
    integer, private :: ndefprop
! this is the index of mobility data, set in init_gtp in subroutine gtp3A
    integer, private :: mqindex
! quasichemical model type, 1=classic, 2=corrcted type 1, 3=corrected type 2
    integer :: qcmodel=1
! this is to remember how manytimes find_gridmeen needs to search all gridp
    integer :: ngridseek

```

8 Subroutines and functions

It is not self evident how to organize the description of the subroutines in GTP. One can base it on the type of service the subroutine provides like entering data, listing data, calculating or on the type of object the action is performed on like on elements, phases, additions etc. A mixed approach is made here where “find”, “enter” and “list” subroutines operating on general objects are grouped together whereas subroutines specific for “state variables”, “additions” and similar things are grouped together for all services like entering, listing and calculations. One reason is that when a new addition is implemented all of these services must be provided together so it is natural to keep them together also in the documentation as a programmer has to provide new subroutines for all of them.

The important part of a subroutine declaration is the arguments and the verbatim section normally include the argument list and a short explanation of those. The local variables and the actual code is not interesting in this documentation. The LaTeX section surrounding the verbatim should explain the action of the subroutine and its connection to the rest of the code. Some functions and subroutines are grouped together in one LaTeX section as they are very similar. But it can be quite a challenge to find a subroutine that perform a desired action. In some cases one may even find two subroutines doing (almost) the same task because I had forgotten or could not find the first version.

8.1 Variable names

The arguments for the subroutines and functions are normally explained but some standards have been used

Symbol	Type	Meaning
iph	integer	Phase index in PHASES
ics	integer	Composition set number
lokph	integer	Phase location (index to PHLISTA)
lokcs	integer	Index of PHASE_VARRES array for a composition set
ceq	pointer	Pointer to current gtp_equilibrium_data record

8.2 Initiallization

This subroutine must be called before any other in the GTP package. It dimensions arrays and creates some initial data structures. The arguments are presently not used but should be used to dimension arrays and provide default values.

```
subroutine init_gtp(intvar,dblvar)
! initiate the data structure
! create element and species record for electrons and vacancies
! the allocation of many arrays should be provided calling this routine
! intvar and dblvar will eventually be used for allocations and defaults
implicit none
```

```
integer intvar(*)
double precision dblvar(*)
```

8.2.1 Initialize of an assessment

This is handled automatically at present.

```
subroutine assessmenthead(ash)
! create an assessment head record and do more (later)
  type(gtp_assessmenthead), pointer :: ash
!   type(gtp_assessmenthead), allocatable :: ash
```

8.3 Functions to know how many and to find them

The counters for elements etc are private so external software must call functions to find out how many elements etc that the system has. They are all given here.

```
integer function noel()
! number of elements because noofel is private
! should take care if elements are suspended
-----
integer function nosp()
! number of species because noofsp is private
! should take care if species are suspended
-----
integer function noph()
! number of phases because noofph is private
! should take care if phases are hidden
-----
integer function noofcs(iph)
! returns the number of compositions sets for phase iph
  implicit none
  integer iph
-----
integer function noconst(iph,ics,ceq)
! number of constituents for iph (include single constituents on a sublattice)
! It tests if a constituent is suspended which can be different in each ics.
  implicit none
  integer iph,ics
  TYPE(gtp_equilibrium_data), pointer :: ceq
-----
integer function nooftup()
! number of phase tuples
```

8.3.1 How many state variable functions

The number of state variable functions entered is given by this.

```
integer function nosvf()  
! number of state variable functions
```

8.3.2 How many equilibria

There is a global array with equilibria records but only a few of them may be allocated with data. This routine returns the number of allocated equilibrium records.

```
integer function noeq()  
! returns the number of equilibria entered
```

8.3.3 Total number of phases and composition sets

This is needed when dimensioning arrays for phases and composition sets for calculations.

```
integer function nonsusphcs(ceq)  
! returns the total number of unhidden phases+composition sets  
! in the system. Used for dimensioning work arrays and in loops  
implicit none  
TYPE(gtp_equilibrium_data), pointer :: ceq
```

8.3.4 Find element, species and components

These subroutines translate from name to index or location of data or vice versa. There are other some “find” subroutines for special things like find_gridmin described in 8.16.5.

```
subroutine find_element_by_name(name,iel)  
! find an element index by its name, exact fit required  
implicit none  
character name*(*)  
integer iel  
-----  
subroutine find_component_by_name(name,icomp,ceq)  
! BEWARE: one may in the future have different components in different  
! equilibria. components are a subset of the species  
implicit none  
character*(*) name  
integer icomp  
TYPE(gtp_equilibrium_data), pointer :: ceq
```

```

-----
subroutine find_species_by_name(name,isp)
! locates a species index from its name, unique abbreviation
! or exact match needed
  implicit none
  character name*(*)
  integer isp
-----

subroutine find_species_record(name,loksp)
! locates a species record allowing abbreviations
  implicit none
  character name*(*)
  integer loksp
-----

subroutine find_species_record_noabbr(name,loksp)
! locates a species record no abbreviations allowed
  implicit none
  character name*(*)
  integer loksp
-----

subroutine find_species_record_exact(name,loksp)
! locates a species record, exact match needed
! for parameters, V must not be accepted as abbreviation of VA or C for CR
  implicit none
  integer loksp
  character name*(*)

```

8.3.5 Find a phase and composition set

These are routines to find a phase by index or name, the phase tuple should be used in application software. Note the phase tuple contain both the phase number and composition set number and also an index to the data structure for the phase.

For the name of the second and higher composition set of a phase use the phase name followed by a hash sign, #, followed by a digit to specify the composition set. If you create composition sets yourself you can add a prefix and suffix to the phase name.

A phase name must not be an abbreviation of the name of another phase. That seems straightforward but sometimes one comes across phase names like ALPHA and ALPHA_PRIME. See also section 8.4.7.

```

subroutine find_phasetuple_by_name(name,phcsx)
! finds a phase with name "name", returns phase tuple index
! handles composition sets either with prefix/suffix or #digit
! When no pre/suffix nor # always return first composition set
  implicit none

```

```

    character name*(*)
    integer phcsx
-----
    subroutine find_phase_by_name(name,iph,ics)
! finds a phase with name "name", returns address of phase, first fit accepted
! handles composition sets either with prefix/suffix or #digit
! When no pre/suffix nor # always return first composition set
    implicit none
    character name*(*)
    integer iph,ics
-----
    integer function find_phasetuple_by_indices(iph,ics)
! subroutine find_phasetuple_by_indices(iph,ics)
! find phase tuple index given phase index and composition set number
    integer iph,ics
-----
    subroutine find_phasesex_by_name(name,phcsx,iph,zcs)
! finds a phase with name "name", returns index and tuplet of phase.
! All phases checked and error return if name is ambiguous
! handles composition sets either with prefix/suffix or #digit or both
! if no # check all composition sets for prefix/suffix
! special if phcsx = -1 and there are several composition sets then
! zcs is set to -(number of composition sets). Used when changing status
! phcsx, iph and zcs are values to return!
    implicit none
    character name*(*)
    integer phcsx,iph,zcs
-----
    subroutine find_phase_by_name_exact(name,iph,ics)
! finds a phase with name "name", returns address of phase. exact match req.
! handles composition sets either with prefix/suffix or #digit
! no pre/suffix nor # gives first composition set
    implicit none
    character name*(*)
    integer iph,ics

```

8.3.6 Find and select equilibrium

As already stated each equilibrium has a separate set of conditions and results. Equilibria are created by calling `enter_equilibrium`, see section 8.6.13. A “default” equilibrium pointed to by the variable “`firsteq`” is created when the program is started. The pointer variable “`ceq`” is used in many subroutine to indicate the “current” equilibrium for which calculations or data are used.

```

    subroutine findeq(name,ieq)
! finds the equilibrium with name "name" and returns its index

```

```

! ieq should be the current equilibrium
  implicit none
  character name*(*)
  integer ieq
-----
  subroutine selecteq(ieq,ceq)
! checks if equilibrium ieq exists and if so set it as current
  implicit none
  TYPE(gtp_equilibrium_data), pointer :: ceq
  integer ieq

```

8.4 Get things

The difference between find and get is not very distinct. Normally the “find” routines requires a name or symbol to return an index or location whereas the “get” routines require an index or location to get more data. Some of the “get” routines are very specific and the documentation is found together with the type of data you want to get.

```

  subroutine get_phase_record(iph,lokph)
! given phase index iph this returns the phase location lokph
  implicit none
  integer iph,lokph
-----
  subroutine get_phase_variance(iph,nv)
! returns the number of independent variable fractions in phase iph
  implicit none
  integer iph,nv
-----
  subroutine get_constituent_location(lokph,cno,loksp)
! returns the location of the species record of a constituent
! required for ionic liquids as phlista is private
  implicit none
  integer lokph,loksp,cno
-----
  subroutine get_phase_compset(iph,ics,lokph,lokcs)
! Given iph and ics the phase and composition set locations are returned
! Checks that ics and ics are not outside bounds.
  implicit none
  integer iph,ics,lokph,lokcs
-----
  subroutine find_constituent(iph,spname,mass,icon)
! find the constituent "spname" of a phase. spname can have a sublattice #digit
! Return the index of the constituent in icon. Additionally the mass
! of the species is returned.
  implicit none

```



```

character*(*) spname
double precision mass
integer iph,icon

```

8.4.1 Get phase constituent name

By supplying a phase index and the sequential index (counted over all sublattices, this routine returns the name and mass of the constituent.

```

subroutine get_constituent_name(iph,iseq,spname,mass)
! find the constituent with sequential index iseq in phase iph
! return name in "spname" and mass in mass
  implicit none
  character*(*) spname
  integer iph,iseq
  double precision mass

```

8.4.2 Get element data

The data for an element is returned.

```

subroutine get_element_data(iel,elsym,elname,refstat,mass,h298,s298)
! return element data as that is stored as private in GTP
  implicit none
  character elsym*2, elname*(*),refstat*(*)
  double precision mass,h298,s298
  integer iel

```

8.4.3 Get component or species name

Components are by default the elements but the user can (sometimes in the future) change this to any set of species (that are orthogonal). Each equilibrium will be able to have a different set of components.

The set of components are important because one can only use components to set amounts of fractions with conditions like $N(< \text{components} >)$. There is an alternative method to set amounts using the subroutine `set_input_amounts` where amounts or mass of different species can be used to give amounts of components.

The `get_species_location` is used when redefining the set of components.

```

subroutine get_component_name(icomp,name,ceq)
! return the name of component icomp
  implicit none
  character*(*) name

```

```

        integer icomp
        TYPE(gtp_equilibrium_data), pointer :: ceq
-----
subroutine get_species_name(isp,spsym)
! return species name, isp is species number
    implicit none
    character spsym*(*)
    integer isp
-----
subroutine get_species_location(isp,loksp,spsym)
! return species location and name, isp is species number
    implicit none
    character spsym*(*)
    integer isp,loksp

```

8.4.4 Get species data relative elements and components

A species is just a stoichiometric arrangement of elements like a molecule. It has no thermodynamic data as they are stored at with the phase where the species is a constituent. The stoichiometry of a species is fixed. The composition of a phase can vary if there are two or more species as constituents in one or more sublattices.

The elements are the simplest species. A species can have a charge which may be non-integer. For some models, like UNIQUAC, species can have additional data used for the modeling like area and volume, see section 8.8.14. These does not constitute a volume for the phase.

The `get_species_component_data` returns the stoichiometry of the species using the current set of components. Normally that the components are the same as the elements.

```

subroutine get_species_data(loksp,nspel,ielno,stoi,smass,qsp,nextra,extra)
! return species data, loksp is from a call to find_species_record
! nspel: integer, number of elements in species
! ielno: integer array, element indices
! stoi: double array, stoichiometric factors
! smass: double, mass of species
! qsp: double, charge of the species
! nextra, integer, number of additional values
! extra: double, some additional values like UNIQUAC volume and area
    implicit none
    integer, dimension(*) :: ielno
    double precision, dimension(*) :: stoi,extra
    integer loksp,nspel,nextra
    double precision smass,qsp
-----
subroutine get_species_component_data(loksp,nspel,compos,stoi,smass,qsp,ceq)
! return species data, loksp is from a call to find_species_record

```

```

! Here we return stoichiometry using components
! nspel: integer, number of components in species
! compno: integer array, component (species) indices
! stoi: double array, stoichiometric factors
! smass: double, mass of species
! qsp: double, charge of the species
  implicit none
  integer, dimension(*) :: compnos
  double precision, dimension(*) :: stoi(*)
  integer loksp,nspel
  double precision smass,qsp
  TYPE(gtp_equilibrium_data), pointer :: ceq

```

8.4.5 New species stoichiometry

A subroutine provided by a user to change the stoichiometry of a species. The reason for this is unclear.

```

subroutine set_new_stoichiometry(loksp, new_stoi, ispel)
! provided by Clement Instoini
! Change the stoichiometric coefficient of the ispel-th element of loksp-th
! species (the last one when ispel is not given)
! loksp: index of the species (input integer)
! new_stoi: new value of the stoichiometric coefficient (input double precision)
! ispel: index of the element (optional, input integer)
  implicit none
  integer, intent(in):: loksp
  integer, intent(in), optional :: ispel
  double precision, intent(in):: new_stoi

```

8.4.6 Mass of component

For the mass balance calculations during equilibrium calculations the mass of a component is needed frequently. This subroutine just returns the mass of a component.

```

double precision function mass_of(component,ceq)
! return mass of component
! smass: double, mass of species
  implicit none
  integer :: component
  TYPE(gtp_equilibrium_data), pointer :: ceq

```

8.4.7 Get phase name

The title says all but there are two variants of this, one using phase tuples, the other integer variables as arguments. There are also some more or less redundant routines to get the record for phase data. See also section 8.3.5.

```
subroutine get_phase_name(iph,ics,name)
! Given the phase index and composition set number this subroutine returns
! the name with pre- and suffix for composition sets added and also
! a \# followed by a digit 2-9 for composition sets higher than 1.
  implicit none
  character name*(*)
  integer iph,ics
-----
subroutine get_phasetup_name(phtupx,name)
! phasetuple(phtupx)%phase is index to phlista
! the name has pre- and suffix for composition sets added and also
! a \# followed by a digit 2-9 for composition sets higher than 1.
  implicit none
  character name*(*)
  integer phtupx
-----
subroutine get_phasetup_name_old(phtuple,name)
! Given the phase tuple this subroutine returns the name with pre- and suffix
! for composition sets added and also a \# followed by a digit 2-9 for
! composition sets higher than 1.
  implicit none
  character name*(*)
  type(gtp_phasetuple) :: phtuple
-----
subroutine get_phasetup_record(phtx,lokcs,ceq)
! return lokcs when phase tuple known
  implicit none
  integer phtx,lokcs
  TYPE(gtp_equilibrium_data), pointer :: ceq
```

8.4.8 Get phase data

This is a very important subroutine used at each iteration during calculations to obtain information about a phase. In the knr array the constituents are given as the integer indices of the species location in the array SPLISTA. The constituents are stored sequentially and the first nkl(1) positions in knr belong to sublattice 1, the next nkl(2) to sublattice 2 etc. The order is alphabetical for each sublattice (not for ionic liquid). In yarr the fractions of the constituents are given sequentially in the same order.

NOTE the ionic liquid has the constituents in the second sublattice ordered by first all anions, then Va (if any), then all neutrals. The anions and the neutrals are ordered alphabetically.

In the array qq the current number of components per formula unit is returned in the first index and the current charge (valence) in the second.

The routine get_phase_structure returns just the number of sublattices and constituents in each. It is needed when calculating the derivatives of the chemical potentials in order to transform mobilities to diffusion coefficients.

```

subroutine get_phase_data(iph,ics,ns1,nkl,knr,yarr,sites,qq,ceq)
! return the structure of phase iph and constitution of comp.set ics
! ns1: integer, number of sublattices
! nkl: integer array, number of constituents in each sublattice
! knr: integer array, species location (not index) of constituents (all subl)
! yarr: double array, fraction of constituents (in all sublattices)
! sites: double array, number of sites in each sublattice
! qq: double array, (must be dimensioned at least 5) although only 2 used:
! qq(1) is number of real atoms per formula unit for current constitution
! qq(2) is net charge of phase for current constitution
! ceq: pointer, to current gtp_equilibrium_data record
  implicit none
  integer, dimension(*) :: nkl,knr
  double precision, dimension(*) :: yarr,sites,qq
  integer iph,ics,ns1
  TYPE(gtp_equilibrium_data), pointer :: ceq
-----
subroutine get_phase_structure(lokph,ns1,nkl)
! return the number of sublattices and constituents in each.
! ns1: integer, number of sublattices
! nkl: integer array, number of constituents in each sublattice
! USED when calculating derivatives of chemical potentials and diffusion coef
  implicit none
  integer, dimension(*) :: nkl
  integer lokph,ns1

```

8.4.9 Phase tuple array

The subroutine get_phtuplearray is redundant as application software can use the global array PHASETUPLE declared within the OC software.

In the phase tuple array the first indices from 1 to nooph represent the first composition set of the phases. Whenever a phase can be stable with two or more compositions extra composition sets are created. These are placed after the first nooph phases.

The function noophphasetuples should be called whenever the grid minimizer has been used as new composition sets may have been created.

```

integer function get_phtuplearray(phcs)
! copies the internal phase tuple array to external software
! function value set to number of tuples
  type(gtp_phasetuple), dimension(*) :: phcs
-----
integer function noofphasetuples()
! number of phase tuples

```

8.5 Set things

Many things can be set but most of the ways to set them are described together with the object to set. How to set conditions is described in 8.8.6 because it involves state variables.

8.5.1 Set constitution

This is a subroutine used frequently when iterating to find the equilibrium. At each iteration the new constitutions of the phases are set using this subroutine. Some internal quantities are also calculated like the number of atoms per mole formula unit of the phase and the charge of the phase. These are returned in the call.

The array yfr in the phase_varres record belonging to the composition set (iph,ics) is not private and a programmer may thus change the values externally without calling set_constitution. But this is strongly discouraged as the internal variables qq(1) and qq(2) must be updated for each set of fractions to ensure that the massbalance is correct.

The order of the fractions in yfra must be the same as in get_phase_data, see 8.4.8.

```

subroutine set_constitution(iph,ics,yfra,qq,ceq)
! set the constituent fractions of a phase and composition set and the
! number of real moles and mass per formula unit of phase
! returns number of real atoms in qq(1), charge in qq(2) and mass in qq(3)
! for ionic liquids sets the number of sites in the sublattices
  implicit none
  double precision, dimension(*) :: yfra,qq
  integer iph,ics
  TYPE(gtp_equilibrium_data), pointer :: ceq

```

8.5.2 Set reference state for a component

For each component the user can select a phase, temperature and pressure as reference state. If a * is given as temperature and pressure the current value of T and P will be used. The reference state is used in calculate_reference_state in section 8.10.12.

```

subroutine set_reference_state(icom,iph,tpval,ceq)
! set the reference state of a component to be "iph" at tpval

```

```

implicit none
integer icomp,iph
double precision, dimension(2) :: tpval
TYPE(gtp_equilibrium_data), pointer :: ceq

```

8.5.3 Set condition

See section 8.8.6.

8.5.4 Amend components

This subroutine changes the set of components of the system. The components are used when setting conditions of the amounts or chemical potentials in the system. By default the elements are the components but in some cases it is convenient to use another set of species for setting conditions. The number of components cannot be changed by this command, the number of new components must be the same as the number of elements. The components must form an orthogonal set.

This command is fragile. Beware that when using other components than the elements one may have negative mole fractions of the components and amounts of the phases (the mass and mass fraction is positive).

```

subroutine amend_components(line,ceq)
! amend the set of components
  implicit none
  character line*(*)
  type(gtp_equilibrium_data), pointer :: ceq

```

8.6 Enter data

The subroutines for entering data and other things can be named as new, enter, add, create etc. according to the mind of the programmer when the subroutine was written. In all cases the data is provided as arguments in the call, there is no interactions with the user.

8.6.1 Enter element data

All data for an element. Some checks are made. The elements are automatically entered also as species so they can be constituents of phases.

```

subroutine enter_element(symb,name,refstate,mass,h298,s298)
! Creates an element record after checks.
! symb: character*2, symbol (it can be a single character like H or V)
! name: character, free text name of the element

```

```

! refstate: character, free text name of reference state.
! mass: double, mass of element in g/mol
! h298: double, enthalpy difference between 0 and 298.14 K
! s298: double, entropy at 298.15 K
  implicit none
  CHARACTER*(*) symb,name,refstate
  DOUBLE PRECISION mass,h298,s298

```

8.6.2 Enter species data

All data for an element. Some checks are made. The elements constituting the species must have been entered before. A species can have a positive or negative charge using the element index -1 with a stoichiometric factor.

```

subroutine enter_species(symb,noelx,ellist,stoik)
! creates a new species
! symb: character*24, name of species, often equal to stoichiometric formula
! noelx: integer, number of elements in stoichiometric formula (incl charge)
! ellist: character array, element names (electron is /-)
! stoik: double array, must be positive except for electron.
  implicit none
  character symb*(*),ellist(*)*(*)
  integer noelx
  double precision stoik(*)

```

8.6.3 Enter phase and model interactively

This routine asks for name and other information needed to enter a phase. Then it calls the routine in next section.

```

subroutine enterphase(ccline,last)
! interactive entering of phase
  character ccline*(*)
  integer last
!   type(gtp_equilibrium_data), pointer :: ceq

```

8.6.4 Enter phase and model

This subroutine is called with the model data needed to create the data structure for a phase (no parameter data). The model variable is just a text, phtype is used to arrange gas (G) and liquids (L) before the alphabetical list of the other phases.

```

subroutine enter_phase(name,ns1,knr,const,sites,model,phtype)

```



```

! creates the data structure for a new phase
! name: character*24, name of phase
! nsl: integer, number of sublattices (range 1-9)
! knr: integer array, number of constituents in each sublattice
! const: character array, constituent (species) names in sequential order
! sites: double array, number of sites on the sublattices
! model: character, free text
! phtype: character*1, specifies G for gas, L for liquid
  implicit none
  character name*(*),model*(*),phtype*(*)
  integer nsl
  integer, dimension(*) :: knr
  double precision, dimension(*) :: sites
  character, dimension(*) :: const*(*)

```

8.6.5 Sorting constituents in ionic liquids

The ionic liquid model requires all cations (with positive charge) on the first sublattice in alphabetical order. On the second sublattice the anions (with negative charge) should be first (in alphabetical order), then the hypothetical vacancy (if any), then any neutrals in alphabetical order. This subroutine takes care of that

```

subroutine sort_ionliqconst(lokph,mode,knr,kconlok,klok)
! sorts constituents in ionic liquid, both when entering phase
! and decoding parameter constituents
! order: 1st sublattice only cations
! 2nd: anions, VA, neutrals
! mode=0 at enter phase, wildcard ok in 1st sublattice if neither anions nor Va
! mode=1 at enter parameter (wildcard allowed, i.e. some kconlok(i)=-1)
! some parameters not allowed, L(ion,A+:B,C), must be L(ion,*:B,C), check!
  implicit none
  integer lokph,knr(*),kconlok(*),klok(*),mode

```

8.6.6 Enter composition set

As explained in section 2.6 a phase may exist simultaneously with several different composition sets. This can be due to miscibility gaps or ordering. Some carbides like cubic TiC is modeled as the same phase as the metallic FCC and it may be stable at the same time as the austenite phase in steels. This subroutine creates a new composition set for a phase.

```

subroutine enter_composition_set(iph,prefix,suffix,icsno)
! adds a composition set to a phase.
! iph: integer, phase index
! prefix: character*4, optional prefix to original phase name

```


8.6.8 Enter parameter

All kind of parameters are entered by this subroutine. Called when reading a TDB file or entered interactively, see 8.8.2.

```
subroutine enter_parameter(lokph,typty,fractyp,nsl,endm,nint,lint,ideg,&
    lfun,refx)
! enter a parameter for a phase from database or interactively
! enter_parameter_inter(activly) is in gtp3D for some unknown reason ...
! typty is the type of property, 1=G, 2=TC, ... , n*100+icon MQ&const#subl
! fractyp is fraction type, 1 is site fractions, 2 disordered fractions
! nsl is number of sublattices
! endm has one constituent index for each sublattice
! constituents in endm and lint should be ordered so endm has lowest
! (done by decode_constarr)
! nint is number of interacting constituents (can be zero)
! lint is array of sublattice+constituent indices for interactions
! ideg is degree
! lfun is link to function (integer index) if -1 used for listing
! refx is reference (text)
! if this is a phase with permutations all interactions should be in
! the first or the first two identical sublattices (except interstitials)
! a value in endm can be negative to indicate wildcard
! for ionic liquid constituents must be sorted specially
    implicit none
    integer, dimension(*) :: endm
    character refx*(*)
    integer lokph,fractyp,typty,nsl,nint,ideg,lfun
    integer, dimension(2,*) :: lint
```

8.6.9 Subroutines handling fcc permutations

These subroutines creates all possible permutations of parameters for a 4 sublattice fcc phase up to ternary interactions. The 4 ordering sublattices must be the first and they represent the tetrahedron in the lattice. The number of sites must be the same and the constituents also. There can be additional sublattices for interstitials. See section 5.6.1.

```
subroutine fccpermut(lokph,nsl,iord,noperm,elinks,nint,jord,intperm,intlinks)
! finds all fcc/hcp permutations needed for this parameter
! The order of elements in the sublattices is irrelevant when one has F or B
! ordering as all permutations are stored in one place (with some exceptions)
! Thus the endmembers are ordered alphabetically in the sublattices and also
! the interaction parameters. Max 2 levels of interactions allowed.
    implicit none
    integer, dimension(*) :: iord,intperm
```

```

integer, dimension(2,*) :: jord
integer lokph,ns1,noperm,nint
-----
subroutine fccip2A(lokph,jord,intperm,intlinks)
! 2nd level interaction permutations for fcc
implicit none
integer, dimension(*) :: intperm
integer, dimension(2,*) :: jord,intlinks
integer lokph
-----
subroutine fccip2B(lq,lokph,lshift,jord,intperm,intlinks)
! 2nd level interaction permutations for fcc
implicit none
integer lq,lokph,lshift
integer, dimension(*) :: intperm
integer, dimension(2,*) :: jord,intlinks
-----
subroutine fccint31(jord,lshift,intperm,intlinks)
! 1st level interaction in sublattice l1 with endmember A:A:A:B or A:B:B:B
! set the sublattice and link to constituent for each endmember permutation
! 1st permutation of endmember: AX:A:A:B; A:AX:A:B; A:A:AX;B 4      0 1 2
! 2nd permutation of endmember: AX:A:B:A; A:AX:B:A; A:A:B:AX 3      0 1 3
! 3rd permutation of endmember: AX:B:A:A; A:B:AX:A; A:B:A:AX 3      0 2 3
! 4th permutation of endmember: B:AX:A:A; B:A:AX:A; B:A:A:AX 1 or 1 2 3
! 1st permutation of endmember: A:BX:B:B; A:B:BX:B; A:B:B:BX 4      0 1 2
! 2nd permutation of endmember: BX:A:B:B; B:A:BX:B; B:A:B:BX 1 etc -1 1 2
! 3rd -1 0 2 ; -1 0 1
! suck
implicit none
integer lshift
integer, dimension(2,*) :: jord,intlinks
integer, dimension(*) :: intperm
-----
subroutine fccint22(jord,lshift,intperm,intlinks)
! 1st level for endmember A:A:B:B with interaction in sublattice jord(1,1)
! 6 permutations of endmember, 2 permutations of interactions, 12 in total
! 1st endmemperm: AX:A:B:B; A:AX:B:B      0 1
! 2nd endmemperm: AX:B:A:B; A:B:AX:B      0 2
! 3rd endmemperm: AX:B:B:A; A:B:B:AX      0 3
! 4th endmemperm: B:AX:B:A; B:A:B:AX      1 3
! 5th endmemperm: B:B:AX:A; B:B:A:AX      2 3
! 6th endmemperm: B:AX:A:B; B:A:AX:B or 1 2
! 1th endmemperm: A:A:BX:B; A:A:B:BX      0 1
! 2nd endmemperm: A:BX:A:B; A:B:A:BX     -1 1
! 3rd endmemperm: A:BX:B:A; A:B:BX:A     -1 0
! 4th endmemperm: BX:A:B:A; B:A:BX:A     -2 0

```

```

! 5th endmemperm: BX:B:A:A; B:BX:A:A      -2 -1
! 6th endmemperm: BX:A:A:B; B:A:A:BX      -2  1
  implicit none
  integer lshift
  integer, dimension(2,*) :: jord,intlinks
  integer, dimension(*) :: intperm
-----
  subroutine fccint211(a211,jord,lshift,intperm,intlinks)
! 1st level interaction in sublattice l1 with endmember like A:A:B:C
! 12 endmember permutations of AABC; ABBC; or ABCC
! 2 interaction permutations for each, 24 in total
  implicit none
  integer a211,lshift
  integer, dimension(2,*) :: jord,intlinks
  integer, dimension(*) :: intperm
-----
  subroutine fccpe211(l1,elinks,nsl,lshift,iord)
! sets appropriate links to constituents for the 12 permutations of
! A:A:B:C (l1=1), A:B:B:C (l1=2) and A:B:C:C (l1=3)
  implicit none
  integer l1,nsl,lshift
  integer, dimension(nsl,*) :: elinks
  integer, dimension(*) :: iord
-----
  subroutine fccpe1111(elinks,nsl,lshift,iord)
! sets appropriate links to 24 permutations when all 4 constituents different
! A:B:C:D
! The do loop keeps the same constituent in first sublattice 6 times, changing
! the other 3 sublattice, then changes the constituent in the first sublattice
! and goes on changing in the other 3 until all configurations done
  implicit none
  integer nsl,lshift
  integer, dimension(nsl,*) :: elinks
  integer, dimension(*) :: iord

```

8.6.10 Subroutines handling bcc permutations

Finally implemented and works!! I hope I never have to look at these subroutines again.

```

  subroutine bccpermut(lokph,nsl,iord,noperm,elinks,nint,jord,intperm,intlinks)
! finds all bcc permutations needed for this parameter
  implicit none
  integer lokph,nsl,noperm,nint
! iord are the endmember constituent indices
! intperm has dimension 24 and contain propagation of interactions ??

```

```

    integer, dimension(*) :: iord,intperm
! jord(1,int) is the interaction subl. and jord(2,int) the constituent index
    integer, dimension(2,*) :: jord
! these must be allocated here and will be stored in the parameter records
! giving the constituent indices for permutations of endmembers and interactions
    integer, dimension(:,:), allocatable :: elinks
    integer, dimension(:,:), allocatable :: intlinks
-----
subroutine bccendmem(lokph,nsl,elal,noperm,elinks)
! generate an bcc endmember with all permutations
    implicit none
    integer lokph,nsl,noperm
! elal are the endmember species indices
    integer, dimension(*) :: elal
! these must be allocated here and will be stored in the parameter records
! giving the sublattice and constituent indices for each permutation
! of an endmembers
    integer, dimension(:,:), allocatable :: elinks
!   integer, dimension(:,:), allocatable :: intlinks
-----
subroutine bccint1(lokph,nsl,elal,noperm,elinks,nint,jord,intperm,intlinks)
! generate all bcc permutations for a first order interaction
    implicit none
! on entry noperm is the number of permutation of the endmember
! on exit  noperm is the number of permutation of the interaction
    integer lokph,nsl,noperm,nint
! elal are the endmember species indices
    integer, dimension(*) :: elal
! intperm has dimension 24 and contain propagation of interactions ??
    integer, dimension(*) :: intperm
! jord(1,int) is the interaction subl. and jord(2,int) the constituent index
    integer, dimension(2,*) :: jord
! these contain the already allocated permutation of the endmember
!   integer, dimension(:,:), allocatable :: elinks
    integer, dimension(nsl,*) :: elinks
! intlinks will be allocated here and will be stored in the parameter records
! giving the constituent indices for permutations of the interactions
! It may be reallocated if the interaction is second level
    integer, dimension(:,:), allocatable :: intlinks
-----
subroutine bccint2(lokph,nsl,elal,noperm,elinks,nint,jord,intperm,intlinks)
! generate all bcc permutations needed for a ternary or reciprocal parameter
    implicit none
! on entry noperm is the number of permutations of the first interaction
! on exit  noperm is the number of permutations of the second interaction
    integer lokph,nsl,noperm,nint

```

```

! elal are the endmember species indices
  integer, dimension(*) :: elal
! intperm has dimension 24 and contain propagation of interactions ??
  integer, dimension(*) :: intperm
! jord(1,int) is the interaction subl. and jord(2,int) the constituent index
  integer, dimension(2,*) :: jord
! elinks are the permutations of the endmember
!   integer, dimension(:,:), allocatable :: elinks
  integer, dimension(nsl,*) :: elinks
! on entry intlinks are the permutations of the first interaction
! on exit  intlinks are the permutations of the second interaction
! it must be deallocated and reallocated using int1links
  integer, dimension(:,:), allocatable :: int1links
  integer, dimension(:,:), allocatable :: intlinks

```

8.6.11 Find constituent

Constituents in a phase are ordered sequentially across all sublattices. This routine finds this sequential index provided with the sublattice number and sequential order in the sublattice.

```

subroutine findconst(lokph,ll,spix,constix)
! locates the constituent index of species with index spix in sublattice ll
! and returns it in constix. For wildcards spix is -99; return -99
! THERE MAY ALREADY BE A SIMILAR SUBROUTINE ... CHECK
  implicit none
  integer lokph,ll,spix,constix

```

8.6.12 Enter the bibliographic reference for parameter data

Bibliographic information for parameters is provided by this subroutine.

```

subroutine tdbrefs(refid,line,mode,iref)
! store a reference from a TDB file or given interactively
! If refid already exist and mode=1 then amend the reference text
  implicit none
  character*(*) refid,line
  integer mode,iref

```

8.6.13 Enter equilibrium

The equilibrium record, as explained in section 6.2.5 has all data necessary for specifying an equilibrium: conditions, compoenets, phases etc. One may have several equilibria with different sets of conditions but they have the same set of phases (the phase status and set of stable phases may differ). This can be used to assess many different experiments or used in simulations where each

gridpoint is connected to an equilibrium record. This simplifies parallel processing as each thread can work independently on the data in the equilibrium record.

```

subroutine enter_equilibrium(name,number)
! creates a new equilibrium. Allocates arrayes for conditions
! components, phase data and results etc.
! returns index to new equilibrium record
! THIS CAN PROBABLY BE SIMPLIFIED, especially phase_varres array can be
! copied as a whole, not each record structure separately ... ???
  implicit none
  character name*(*)
  integer number
-----
subroutine geneqname(text)
! creates a equilibrium name like EQ_x where x is the freeq in text
  implicit none
  character text*(*)

```

8.6.14 Enter many equilibria

This command is specially designed for entering table of experimental data where most conditions are the same but only some values are different. It will ask for a table head which must contain all information needed to calculate the equilibrium. By default all phases are suspended so first give the set of phases to be considered (entered, fixed or dormant). The line with the phase status start with the status, for fix and entered followed by a number and then a list of phases. An asterisk, *, can be used for all phases.

The conditions can be set after just the word “conditions” and experiments after the word “experiment”. It is also possible to set reference states and to demand that some symbols are calculated or values of state variables and parameter identifiers listed.

Each line may refer to columns in the table to follow after the head. The columns are specified with teh “@” character followed by the column. Max 9 columns allowed.

The head is finished by the command “table_start” and then on each line the column values must be given. But first on each line there must be a name of the equilibrium (this is not counted as a column, or as column zero).

The lines with values is terminated by a “table_end”.

All equilibria in a table can be calculated with the “calculate all” command after the range of equilibria has been set by the “set range” command.

```

subroutine enter_many_equil(cline,last,pun)
! executes an enter many_equilibria command
! and creates many similar equilibria from a table
! pun is file units for storing experimental dataset, pun(i)>0 if i is open
  implicit none

```



```

character*(*) cline
integer last,pun(9)

```

8.6.15 Delete all conditions

When amending the set of components all conditions are also deleted for all equilibria. This subroutine is also used when one gives the command SET CONDITION *:=NONE. To avoid memory leaks it is also used when deleting an equilibrium.

```

subroutine delete_all_conditions(mode,ceq)
! deletes the (circular) list of conditions in an equilibrium
! it also deletes any experiments
! if mode=1 the whole equilibrium is removed, do not change phase status
! because the phase_varres records have been deallocated !!!
! I am not sure it releases any memory though ...
  implicit none
  integer mode
  type(gtp_equilibrium_data), pointer :: ceq

```

8.6.16 Delete equilibria

This is needed after STEP or MAP to clean up the structure as all equilibria along the lines are saved as equilibrium records.

```

subroutine delete_equilibria(name,ceq)
! deletes equilibria (needed when repeated step/map)
! name can be an abbreviation line "_MAP*"
! deallocates all data. Minimal checks ... one cannot delete "ceq"
  implicit none
  character name*(*)
  type(gtp_equilibrium_data), pointer :: ceq

```

8.6.17 Copy equilibrium

As part of STEP and MAP equilibrium records are copied between different lists. The second variant also specifies the index of the equilibrium in the eqlista array.

```

subroutine copy_equilibrium(neweq,name,ceq)
! creates a new equilibrium which is a copy of ceq.
  implicit none
  character name*(*)
  type(gtp_equilibrium_data), pointer :: neweq,ceq
-----

```

```

subroutine copy_equilibrium2(neweq,number,name,ceq)
! creates a new equilibrium which is a copy of ceq. THIS IS STILL USED !! ??
! Allocates arrayes for conditions,
! components, phase data and results etc. from equilibrium ceq
! returns a pointer to the new equilibrium record
! THIS CAN PROBABLY BE SIMPLIFIED, especially phase_varres array can be
! copied as a whole, not each record structure separately ... ???
  implicit none
  character name*(*)
  integer number
  type(gtp_equilibrium_data), pointer ::neweq,ceq

```

8.6.18 Copy condition

This is a utility used in MAP and STEP

```

subroutine copy_condition(newrec,oldrec)
! Creates a copy of the condition record "oldrec" and returns a link
! to the copy in newrec. The links to "next/previous" are nullified
  implicit none
  type(gtp_condition), pointer :: oldrec
  type(gtp_condition), pointer :: newrec

```

8.6.19 Check that a phase is allowed to have fcc permutations

Some minimal checks made when the user or TDB tries to set that a phase can have permutations. Only for 4 sublattice ordering in FCC.

```

logical function check_minimal_ford(lokph)
! some tests if the fcc/bcc permutation model can be applied to this phase
! The function returns FALSE if the user may set the FORD or BORD bit of lokph
  implicit none
  integer lokph

```

8.6.20 Calculate new highcs value when a composition set is created or deleted

Highcs is used when saving the allocated phase_varres records and in some other cases. Note that there can be unused phase_varres records below highcs. The handling of resrving and releasing records in the phase_varres array is not well implemented but it seems to work.

```

integer function newhighcs(reserved)
! updates highcs and arranges csfree to be in sequential order
! highcs is the highest used varres record before the last reservation

```

```

! or release of a record.  release is TRUE if a record has been released
! csfree is the beginning of the free list of varres records.
  implicit none
  logical reserved

```

8.7 List things

The routines in this section are intended for the line oriented user interface of GTP. It lists data assuming 80 column width of the screen. In some cases a character variable is returned but in most case the list unit is provided in the call. This can be the screen or a file.

Some listings are described in connection with the objects that are listed, see 8.14.13.

8.7.1 List data for all elements

The element data is listed. Second version for TDB files.

```

subroutine list_all_elements(unit)
! lists elements
  implicit none
  integer unit
-----
subroutine list_all_elements2(unit)
! lists elements
  implicit none
  integer unit

```

8.7.2 List data for all components

The components may be different in each equilibrium.

```

subroutine list_all_components(unit,ceq)
! lists the components for an equilibrium
  implicit none
  integer unit
  TYPE(gtp_equilibrium_data), pointer :: ceq

```

8.7.3 List data for one element

The data for element “elno” is written to the character variable text from position ipos.

```

subroutine list_element_data(text,ipos,elno)
  implicit none
  character text*(*)
  integer ipos,elno

```

8.7.4 List data for one species

The data for species “spno” is written to the character variable text from position ipos. The second version is suitable for TDB files.

```
subroutine list_species_data(text,ipos,spno)
  implicit none
  character text*(*)
  integer ipos,spno
  -----
subroutine list_species_data2(text,ipos,loksp)
! loksp is species record ...
  implicit none
  character text*(*)
  integer ipos,loksp
```

8.7.5 List data for all species

One line for each species is listed on device unit.

```
subroutine list_all_species(unit)
  implicit none
  integer unit
```

8.7.6 List sorted phases

The phases are listed with one line each, first the stable phases, then the entered but unstable in decreasing order of stability. If more than 10 phases the remaining are merged into a single line.

```
subroutine list_sorted_phases(unit,ceq)
! short list with one line for each phase
! suspended phases merged into one line
! stable first, then entered ordered in driving force order, then dormant
! also in driving force order. Only 10 of each, the others lumped together
  implicit none
  integer unit
  TYPE(gtp_equilibrium_data), pointer :: ceq
```

8.7.7 List a little data for all phases

One line for each phase is listed on device unit for equilibrium ceq.

```
subroutine list_all_phases(unit,ceq)
```

```

! short list with one line for each phase
! suspended phases merged into one line
  implicit none
  integer unit
  TYPE(gtp_equilibrium_data), pointer :: ceq

```

8.7.8 List global results

This is part of the “list_result” command in the GTP user i/f.

```

subroutine list_global_results(lut,ceq)
! list G, T, P, V and some other things
  implicit none
  integer lut
  TYPE(gtp_equilibrium_data), pointer :: ceq

```

8.7.9 List components result

This is part of the “list_result” command in the GTP user i/f.

```

subroutine list_components_result(lut,mode,ceq)
! list one line per component (name, moles, x/w-frac, chem.pot. reference state
! mode 1=mole fractions, 2=mass fractions
  implicit none
  integer lut,mode
  TYPE(gtp_equilibrium_data), pointer :: ceq

```

8.7.10 List all phases with positive dgm

This is part of the “list_result” command in the GTP user i/f. If a phase has positive DGM it should either be dormant or there has been an error calculating the equilibrium.

```

subroutine list_phases_with_positive_dgm(mode,lut,ceq)
! list one line for each phase+comp.set with positive dgm on device lut
! The phases must be dormant or the result is in error. mode is not used
  implicit none
  integer mode,lut
  TYPE(gtp_equilibrium_data), pointer :: ceq

```

8.7.11 List results for one phase

This is part of the “list_result” command in the GTP user i/f. It lists normally only the stable phases with their amounts and compositions. With different values of mode the listing can be changed. Rather clumsy.

```

subroutine list_phase_results(iph,jcs,mode,lut,once,ceq)
! list results for a phase+comp.set on lut
! mode specifies the type and amount of results,
! unit digit: 0=mole fraction,      othewise mass fractions
! 10th digit: 0=only composition,   10=also constitution
! 100th digit: 0=value order,       100=alphabetical order
! 1000th digit: 0=all phases,       1000=only stable phases
! 10000th digit: 1= constituent fractions times formula unit of phase (Solgas)
implicit none
integer iph,jcs,mode,lut
logical once
TYPE(gtp_equilibrium_data), pointer :: ceq
-----
subroutine list_short_results(lut,ceq)
! list short results for all stable phases (for debugging) lut
implicit none
integer lut
TYPE(gtp_equilibrium_data), pointer :: ceq

```

8.7.12 Formatted output for constitution

This subroutine formats the output of composition or constitution in nice columns trying to use as few lines as possible.

```

subroutine format_phase_composition(mode,nv,consts,vals,lut)
! list composition/constitution in alphabetical or value order
! entalsiffra 0 mole fraction, 1 mass fraction, 3 mole percent, 4 mass percent
! tiotalsiffra alphabetical order ... ??
! mode >100 else alphabetical order
! nv is number of components/constitunents (in alphabetical order in consts)
! components/constituents in consts, fractions in vals
implicit none
integer nv,mode,lut
character consts(nv)*(*)
double precision vals(nv)

```

8.7.13 List data on SCREEN or TDB, LaTeX or macro format

This subroutines list the model parameters for all phases in a selected format. Only SCREEN and TDB are implemented.

```

subroutine list_many_formats(cline,last,ftyp,unit1)
! lists all data in different formats: SCREEN/TDB/MACRO/LaTeX/ODB
!
!
!
!
!

```

```

! unfinished
  implicit none
  character cline*(*)
  integer last,unit1,ftyp

```

8.7.14 List some phase model stuff

This is probably redundant but can be used to check the conversion from site fractions to disordered fractions for phases with several fraction sets.

```

subroutine list_phase_model(iph,ics,lut,CHTD,ceq)
! list model (no parameters) for a phase on lut
  implicit none
  integer iph,ics,lut
  TYPE(gtp_equilibrium_data), pointer :: ceq
  character CHTD*1

```

8.7.15 List all parameter data for a phase

This is the big listing of the model and data for a phase. It lists the sublattices, sites, constituents. Then all endmembers and all interaction parameters.

The second version is suitable for TDB files.

```

subroutine list_phase_data(iph,CHTD,lut)
! list parameter data for a phase on unit lut
  implicit none
  integer iph,lut
  character CHTD*1
-----
subroutine list_phase_data2(iph,ftyp,CHTD,lut)
! list parameter data for a phase on unit lut in ftyp format, ftyp=2 is TDB
  implicit none
  integer iph,lut,ftyp
  character CHTD*1

```

8.7.16 Format expression of references for endmembers

When listing an endmember parameter for the Gibbs energy this subroutine subtracts the H298 expression.

```

subroutine subrefstates(funexpr,jp,lokph,parlist,endm,noelin1)
! list a sum of reference states for a G parameter
! like "-H298(BCC_A2,FE)-3*H298(GRAPITE,C)"

```

```

implicit none
integer jp,lokph,parlist,endm(*)
character funexpr*(*)
logical noelin1

```

8.7.17 Encode stoichiometry of species

This subroutine generate a stoichiometric formula for a species including a charge.

```

subroutine encode_stoik(text,ipos,spno)
! generate a stoichiometric formula of species from element list
implicit none
integer ipos,spno
character text*(*)

```

8.7.18 Decode stoichiometry of species

This subroutine can translate a stoichiometric formula to elements and stoichiometric factors including a charge.

```

subroutine decode_stoik(name,noelx,elsyms,stoik)
! decode a species stoichiometry in name to element index and stoichiometry
! all in upper case
implicit none
character name*(*),elsyms(*)*2
double precision stoik(*)
integer noelx

```

8.7.19 Encode constituent array for parameters

This subroutine generates a constituent array for a parameter. Constituents are species. Constituents in different sublattices are separated by “:”, interacting constituents in same sublattice are separated by “,”. The degree is written after a “,”.

```

subroutine encode_constarr(constarr,ns1,endm,nint,lint,ideg)
! creates a constituent array
implicit none
character constarr*(*)
integer, dimension(*) :: endm
integer ns1,nint,ideg
integer, dimension(2,*) :: lint

```


8.7.20 Decode constituent array for parameters

By providing the indices of constituents in the endmember and possible interaction constituents and the degree, a text with the constituent array is generated.

```
subroutine decode_constarr(lokph,constarr,ns1,endm,nint,lint,ideg)
! deconde a text string with a constituent array
! a constituent array has <species> separated by , or : and ; before degree
  implicit none
  character constarr*(*)
  integer endm(*),lint(2,*)
  integer ns1,nint,ideg,lokph,lord
```

8.7.21 List parameter data references

This subroutine lists the source of one or several bibliographic references for the parameters.

```
subroutine list_bibliography(bibid,lut)
! list bibliographic references
  implicit none
  integer lut
  character bibid*(*)
```

8.7.22 List conditions on a file or screen

The heading says all. After the list the degrees of freedom is shown.

```
subroutine list_conditions(lut,ceq)
! lists conditions on lut
  implicit none
  integer lut
  type(gtp_equilibrium_data), pointer :: ceq
```

8.7.23 Extract one condition or experiment in a character variable

A single condition is written in a character variable. The get_one_experiment routine returns the a text line describing an experiment.

```
subroutine get_one_condition(ip,text,seqz,ceq)
! list the condition with the index seqz into text
! It lists also fix phases and conditions that are not active
  implicit none
  integer ip,seqz
```

```

character text*(*)
TYPE(gtp_equilibrium_data), pointer :: ceq
-----
subroutine get_one_experiment(ip,text,seqz,eval,ceq)
! list the experiment with the index seqz into text
! It lists also experiments that are not active ??
! UNFINISHED current value should be appended
implicit none
integer ip,seqz
character text*(*)
logical eval
TYPE(gtp_equilibrium_data), pointer :: ceq

```

8.7.24 List condition in character variable

All current active conditions in equilibrium ceq is written to the character variable text. This can be written on the screen or used for other purposes. It can also be used for experiments (not implemented yet).

```

subroutine get_all_conditions(text,mode,ceq)
! list all conditions if mode=0, experiments if mode=1, -1 if no numbers
implicit none
integer mode
character text*(*)
TYPE(gtp_equilibrium_data), pointer :: ceq

```

8.7.25 Get the degrees of freedom

The current number of degrees of freedom. To calculate an equilibrium it must be zero.

```

integer function degrees_of_freedom(ceq)
! returns the degrees of freedom
implicit none
TYPE(gtp_equilibrium_data), pointer :: ceq

```

8.7.26 List available parameter identifiers

The GTP package allows definition of new properties that can be modeled as dependent on the constitution of each phase. Such properties must be defined in the software and they can be listed with this subroutine. Many additions depend on such parameter properties like the Curie temperature and the Debye temperature.

One can also add properties that does not affect the Gibbs energy but which depend on the constitution of the phase like the mobility, resistivity, lattice parameter etc.

```

subroutine list_defined_properties(lut)
! lists all parameter identifiers allowed
  implicit none
  integer lut

```

8.7.27 Find defined properties

Although properties like TC (Curie temperature) and BMAG (Average Bohr magneton number) are not state variables they can be listed using the command LIST STATE_VARIABLES and their values can be obtained by the same subroutines that are used for state variables like get_state_variable. They use the following subroutine to find the properties defined in the gtp_propid structure.

```

subroutine find_defined_property(symbol,mode,typty,iph,ics)
! searches the propid list for one with symbol or identification typty
! if mode=0 then symbol given, if mode=1 then typty given
! symbol can be TC(BCC), BM(FCC), MQ&FE(HCP) etc, the phase must be
! given in symbol as otherwise it is impossible to find the consititent!!!
! A constituent may have a sublattice specifier, MQ&FE#3(SIGMA)
  implicit none
  integer mode,typty,iph,ics
  character symbol*(*)

```

8.7.28 List some odd details

Creating a line with all phase that are stable (dgm=0). It is used when listing step/map results.

```

subroutine line_with_phases_withdgm0(line,ceq)
! used in amend lines with stored STEP/MAP results
! enter first 6, two .. and last 2 characters of phase names with abs(dgm)<1-8
! line LIQUID#2, PHYRRO..#2
  implicit none
  TYPE(gtp_equilibrium_data), pointer :: ceq
  character line*(*)

```

8.7.29 List more odd details

I do not remember what this is used for.

```

subroutine list_equilibria_details(mode,teq)
! not used yet ... ??
  implicit none
  TYPE(gtp_equilibrium_data), pointer :: teq
  integer mode

```

8.7.30 List an error message if any

The error code is a global variable `gxanytime`. If not zero there has been an error detected and usually there is a message associated with the value. This routine can list the message.

```
logical function gtp_error_message(reset)
! tests the error code and writes the error message (if any)
! and reset error code if reset=0
! if reset >0 that is set as new error message
! if reset <0 the error code is not changed
! return TRUE if error code set, FALSE if error code is zero
  implicit none
  integer reset
```

8.7.31 List coefficients used in an assessment

Listing coefficients used in assessment of model parameters.

```
subroutine listoptcoeff(mexp,error2,done,lut)
! listing of optimizing coefficients
  integer lut,mexp
! error2 is an array with 1: old error, 2: new error; 3: normalized error
  double precision error2(*)
  logical done
!   integer lut,mexp
!   double precision errs(*)
!   type(gtp_equilibrium_data), pointer :: ceq
```

8.8 Interactive subroutines

The current user interface to OC and GTP is command oriented and there are subroutines provided in GTP to enter, set, list and get many things. Most subroutines where the user is expected to provide information is collected in this section.

8.8.1 Ask for phase constitution

The user can interactively set the default constitution or enter a constitution for a specific phase and composition set.

```
subroutine ask_phase_constitution(cline,last,iph,ics,lokcs,ceq)
! interactive input of a constitution of phase iph
  implicit none
  integer last,iph,ics,lokcs
```

```

character cline*(*)
-----
subroutine ask_phase_new_constitution(cline,last,iph,ics,lokcs,ceq)
! interactive input of a constitution of phase iph
  implicit none
  integer last,iph,ics,lokcs
  character cline*(*)

```

8.8.2 Ask for parameter

The user can enter a model parameter with this subroutine. The routine that enters the parameter in the data structure is described in section 8.6.8.

```

subroutine enter_parameter_interactivly(cline,ip,mode)
! enter a parameter from terminal or macro
! NOTE both for ordered and disordered fraction set !!
! mode = 0 for entering
!       1 for listing on screen (kou)
  implicit none
  integer ip,mode
  character cline*(*)

```

8.8.3 Amend global bits

There are a number of global bits that can be set by this subroutine.

```

subroutine amend_global_data(cline,ipos)
  implicit none
  character cline*(*)
  integer ipos

```

8.8.4 Ask for reference of parameter data

Each parameter in a model should have a bibliographic reference, preferably a published paper. When a parameter is entered by calling `enter_parameter_interactivly` the reference is asked for but with this routine it is possible to enter such a reference separately.

```

subroutine enter_bibliography_interactivly(cline,last,mode,iref)
! enter a reference for a parameter interactivly
! mode=0 means enter, =1 amend
  implicit none
  character cline*(*)
  integer last,mode,iref
  logical twotries

```

8.8.5 Enter an experiment

With this subroutine the state variable for an experiment, its value and uncertainty is given for equilibrium ceq. The experiment can also be changed or removed (set equal to NONE).

The logical function is used to check if two state variable records represent the same state variable (because associated would only be true if they were the same record).

```
subroutine enter_experiment(cline,ip,ceq)
! enters an experiment, almost the same as set_condition
  implicit none
  character cline*(*)
  integer ip
  type(gtp_equilibrium_data), pointer :: ceq
-----
  logical function same_statevariable(svr1,svr2)
! returns TRUE if the state variable records are identical
  type(gtp_state_variable), pointer :: svr1,svr2
```

8.8.6 Set a condition

This is the central routine to set a condition for an equilibrium calculation. An alternative is the set_input_amount. When setting the status of a phase as fixed this subroutine is called automatically to add this as condition.

```
subroutine set_condition(cline,ip,ceq)
! to set a condition
  implicit none
  character cline*(*)
  integer ip
  type(gtp_equilibrium_data), pointer :: ceq
-----
  subroutine set_cond_or_exp(cline,ip,new,notcond,ceq)
! decode an equilibrium condition, can be an expression with + and -
! the expression should be terminated with an = or value supplied on next line
! like "T=1000", "x(liq,s)-x(pyrrh,s)=0", "mu(cr)-1.5*mu(o)=muval"
! Illegal with number before first state variable !!!
! It can also be a "NOFIX=<phase>" or "FIX=<phase> value"
! The routine should also accept conditions identified with the "<number>:"
! where <number> is that preceeding each condition in a list_condition
! It should also accept changing conditions by <number>:=new_value
! The pointer to the (most recent) condition or experiment is returned in new
! notcond is 0 if a condition should be created, otherwise an experiment
  implicit none
  integer ip,notcond
```

```

character cline*(*)
TYPE(gtp_equilibrium_data), pointer :: ceq
TYPE(gtp_condition), pointer :: new
-----
subroutine get_experiment_with_symbol(symsym,experimenttype,temp)
! finds an experiment with s symbol index symsym and exp.type
  implicit none
  integer symsym,experimenttype
  type(gtp_condition), pointer :: temp
! NOTE: temp must have been set to ceq%lastcondition before calling this

```

8.8.7 Get condition record

A condition is typically a state variable assigned a value like $T = 1273$ but it can be much more complicated. One can have conditions like $x(\text{liquid}, S) - x(\text{pyrrhotite}, S) = 0$ to specify the congruent melting point of pyrrhotite. A condition can also be that a phase is fixed, i.e. prescribed to be stable.

A condition is specified by the number of terms, the state variable with possible indices (to specify phase, component or constituent etc), reference state (for chemical potentials) and unit (Joule or calorie or per cent or fraction)

The state variable record was not used when this subroutine was first written and the original version is now called `get_condition2`. This version is depreciated and should not be used. The subroutine that replaces this has the original name and has a state variable record as argument. An additional subroutine that returns the state variable record is also available.

The first two subroutines return a pointer to the condition record, see `gtp_condition` for that structure, or an error code if no such condition.

The subroutine to set conditions is described in 8.8.6.

```

subroutine get_condition(nterm,svr,pcond)
! finds a condition/experiment record with the given state variable expression
! If nterm<0 svr is irrelevant, the absolute value of nterm is the sequential
! number of the ACTIVE conditions
  implicit none
  integer nterm
  type(gtp_state_variable), pointer :: svr
! NOTE: pcond must have been set to ceq%lastcondition before calling this
! pcond: pointer, to a gtp_condition record for this equilibrium
  type(gtp_condition), pointer :: pcond
-----
subroutine get_condition2(nterm,coeffs,istv,indices,iref,iunit,pcond)
! finds a condition record with the given state variable expression
! nterm: integer, number of terms in the condition expression
! istv: integer, state variable used in the condition

```

```

! indices: 2D integer array, state variable indices used in the condition
! iref: integer, reference state of the condition (if applicable)
! iunit: integer, unit of the condition value
! NOTE: pcond must have been set to ceq%lastcond before calling this routine!!!
! pcond: pointer, to a gtp_condition record for this equilibrium
! NOTE: conditions like expressions  $x(\text{mg}) - 2 \cdot x(\text{si}) = 0$  not implemented
! fix phases as conditions have negative condition variable
  implicit none
  TYPE(gtp_condition), pointer :: pcond
  integer, dimension(4,*) :: indices
  integer nterm,istv,iref,iunit
  double precision coeffs(*)
-----
  subroutine extract_stvr_of_condition(pcond,nterm)
! finds a condition record with the given state variable record
! returns it as a state variable record !!!
! nterm: integer, number of terms in the condition expression
! pcond: pointer, to a gtp_condition record
  implicit none
  TYPE(gtp_condition), pointer :: pcond
  integer nterm

```

8.8.8 A utility routine to locate a condition record

This is needed by the STEP and MAP commands to find a condition to use as axis variable.

```

  subroutine locate_condition(seqz,pcond,ceq)
! locate a condition using a sequential number
  implicit none
  integer seqz
  type(gtp_condition), pointer :: pcond
  type(gtp_equilibrium_data), pointer :: ceq

```

8.8.9 A utility routine to get the current value of a condition

This subroutine is called several times at each iteration in the equilibrium calculation to formulate the system matrix containing the external conditions. It is still very rudimentary and can be improved.

We do not need to pass the pointer to ceq as that is found via the condition record?

```

  subroutine apply_condition_value(current,what,value,cmix,ccf,ceq)
! This is called when calculating an equilibrium.
! It returns a condition at each call, at first call current must be nullified?
! When all conditions done the current is nullified again

```



```

! If what=-1 then return degrees of freedoms and maybe something more
! what=0 means calculate current values of conditions
! calculate the value of a condition, used in minimizing G
! ccf are the coefficients for conditions with several terms
  implicit none
  integer what,cmix(*)
  double precision value,ccf(*)
  TYPE(gtp_equilibrium_data), pointer :: ceq
  TYPE(gtp_condition), pointer :: current
-----
  subroutine condition_value(mode,pcond,value,ceq)
! set (mode=0) or get (mode=1) a new value of a condition.  Used in mapping
  implicit none
  integer mode
  type(gtp_condition), pointer :: pcond
  type(gtp_equilibrium_data), pointer :: ceq
  double precision value

```

8.8.10 Ask for default phase constitution

The user can enter a default constitution for a phase. A negative value means a maximum value, a positive means a minimum value.

```

  subroutine ask_default_constitution(ccline,last,iph,ics,ceq)
! set values of default constitution interactively
! phase and composition set already given
  implicit none
  character ccline*(*)
  integer last,iph,ics
  TYPE(gtp_equilibrium_data), pointer :: ceq

```

8.8.11 Set default constitution

The user can enter a “default” constitution of a phase. This can sometimes be used as start values for a calculation.

```

  subroutine enter_default_constitution(iph,ics,mmyfr,ceq)
! user specification of default constitution for a composition set
  implicit none
  TYPE(gtp_equilibrium_data), pointer :: ceq
  integer iph,ics
  real mmyfr(*)

```

8.8.12 Interactive set input amounts

This subroutine allows setting condition by entering the amount of species. The amount of the species is converted to amount of components internally. Redundancy is allowed. If several species contain the same element the amounts are added. For example set_input_amount N(C1O1)=10, N(H2O1)=5, N(C1H4)=7, N(O2)=20 is translated to the conditions N(C)=17, N(H)=38, N(O)=55.

```
subroutine set_input_amounts(cline,lpos,ceq)
! set amounts like n(specie)=value or b(specie)=value
! value can be negative removing amounts
! values are converted to moles and set or added to conditions
  implicit none
  integer lpos
  character cline*(*)
  TYPE(gtp_equilibrium_data), pointer :: ceq
```

8.8.13 Utility to decode a model parameter identifier

Used when entering data. The model parameter identifiers are listed in Table 1.

```
subroutine get_parameter_tyty(name1,lokph,tyty,fractyp)
! interpret parameter identifiers like MQ&C#2 in MQ&C#2(FCC_A1,FE:C) ...
! find the property associated with this symbol
  implicit none
  integer tyty,fractyp,lokph
  character name1*(*)
```

8.8.14 Entering a species property

This is used for adding some property to a species, for example the value and area for a species in the UNIQUAC model.

```
subroutine enter_species_property(loksp,nsp, value)
! enter an extra species property for species loksp
  implicit none
  integer loksp,nsp
  double precision value
-----
subroutine set_uniquac_species(loksp)
! set the status bit and allocates spexttra array
  implicit none
  integer loksp
```

8.8.15 Enter material

This is a tentative routine to enter a material rather than reading directly a database.

```
subroutine enter_material(cline,last,nv,xknown,ceq)
! enter a material from a database
! called from user i/f
  implicit none
  integer last,nv
  character cline*(*)
  double precision xknown(*)
  type(gtp_equilibrium_data), pointer :: ceq
```

8.9 Save and read data from files

Most of the subroutines in this section are unfinished. The only thing that can be read is a simple TDB file.

The problem is that whenever a change is made in the data structure these routines must be modified accordingly. And the data structure is still undergoing big changes.

These subroutines are in total disorder

8.9.1 Save all data

Not implemented yet.

```
subroutine gtpsave(filename,str)
! save all data on file, unformatted, TDB or macro
! header
! element list
! species list
! phase list with sublattices, endmembers, interactions and parameters etc
! tpfuns
! state variable functions
! references
!
  implicit none
  character*(*) filename,str
```

8.9.2 Save data in LaTeX and other formats

This is just a dream.

```

subroutine gtpsavelatex(filename,specification)
! save all data on LaTeX format on a file (for publishing)
! header
! element list
! species list
! phase list with sublattices, endmembers, interactions and parameters etc
! tpfuns
! state variable functions
! references
! equilibrium record(s) with conditions, componenets, phase_varres records etc
! anything else?
  implicit none
  character*(*) filename,specification
-----
subroutine gtpsavedir(filename,specification)
! save all data on a direct file (random access)
! header
! element list
! species list
! phase list with sublattices, endmembers, interactions and parameters etc
! tpfuns
! state variable functions
! references
! equilibrium record(s) with conditions, componenets, phase_varres records etc
! anything else?
  implicit none
  character*(*) filename,specification
-----
subroutine gtpsavetm(filename,str)
! save all data on file in macro format
! header
! element list
! species list
! phase list with sublattices, endmembers, interactions and parameters etc
! tpfuns
! state variable functions
! references
!
  implicit none
  character*(*) filename,str
-----
subroutine gtpsavetdb(filename,specification)
! save all data in TDB format on an file UNFINISHED
! header
! element list
! species list

```

```

! phase list with sublattices, endmembers, interactions and parameters etc
! tpfuns
! state variable functions
! references
! equilibrium record(s) with conditions, componenets, phase_varres records etc
! anything else?
  implicit none
  character*(*) filename,specification

```

8.9.3 Save all data unformatted on a file

This is fragile but seems to work. All data are converted to an integer array where pointers are converted to indices in this array. That is the old save format used in Thermo-Calc. Then this array is written unformatted on a file. It can be read back and all data restored.

There is no way to save STEP or MAP results this way as that need a direct (random access) file but experimental data can be saved. Thus it is similar to a PAR file in Thermo-Calc.

It calls a large number of routines included list below.

```

subroutine gtpsaveu(filename,specification)
! save all data unformatted on an file
! First move it to an integer workspace, then write that on a file
! header
! element list
! species list
! phase list with sublattices, endmembers, interactions and parameters etc
! tpfuns
! state variable functions
! references
! equilibrium record(s) with conditions, componenets, phase_varres records etc
! anything else?
  implicit none
  character*(*) filename,specification
-----
subroutine savephases(phroot,iws)
! save data for all phases and store pointer in iws(phroot)
! For phases with disordered set of parameters we must access the number of
! sublattices via firsteq
  implicit none
  integer phroot,iws(*)
-----
subroutine saveequil(lok1,iws)
! subroutine saveequil(lok1,iws,ceq)
! save data for equilibrium record ceq including phase_varres
  implicit none

```

```

    integer lok1,iws(*),jeq
-----
    subroutine svfunsave(loksvf,iws,ceq)
! saves all state variable functions as texts in iws
    implicit none
    integer iws(*),loksvf
    type(gtp_equilibrium_data), pointer :: ceq
-----
    subroutine bibliosave(bibhead,iws)
! saves references on a file
    implicit none
    integer bibhead,iws(*)
-----
    subroutine saveash(lok,iws)
! saving assessment records
    integer lok,iws(*)
-----
    integer function ceqrecsize()
! calculates the number of words needed to save an equilibrium record

```

8.9.4 Read unformatted data saved on file

This reverses the action of save unformatted. The OC data structure is filled with data earlier saved on a file. It is fragile.

```

    subroutine gtpread(filename,str)
! read unformatted all data in the following order
! header
! element list
! species list
! phase list with sublattices, endmembers, interactions and parameters etc
! tpfuns
! references
! first equilibrium record with conditions, componenets, phase_varres etc
! state variable functions
! equilibrium record(s) with conditions, componenets, phase_varres, experim etc
! CCI changed to use iso_fortran_env to find file unit number for C++
    use :: iso_fortran_env
! CCI
    implicit none
    character*(*) filename,str
-----
    subroutine readphases(kkk,iws)
! read data for phlista and all endmembers etc
! works for test case without disordered fraction test

```

```

    implicit none
    integer kkk,iws(*)
-----
    subroutine readendmem(lokem,iws,nsl,emrec)
! allocates and reads an endmember record and its property record from iws
! emrec is an un-allocated pointer in the parameter tree structure
    implicit none
    integer lokem,nsl,iws(*)
    type(gtp_endmember), pointer :: emrec
-----
    subroutine readproprec(lokpty,iws,firstproprec)
! allocates and a property record for both endmembers and interactions
    implicit none
    integer lokpty,iws(*)
    type(gtp_property), pointer :: firstproprec
-----
    subroutine readintrec(lokint,iws,level,intrec)
! allocates and reads an interaction record
    implicit none
    integer lokint,iws(*),level
    type(gtp_interaction), pointer :: intrec
-----
    subroutine readequil(lokeq,iws,elope)
! subroutine readequil(lokeq,iws,elope,ceq)
! lokeq is index for equilibrium record in iws
    implicit none
    integer lokeq,iws(*),elope
-----
    subroutine svfunread(loksvf,iws)
! read a state variable function from save file and store it.
! by default there are some state variable functions, make sure
! they are deleted. Done here just by setting nsvfun=0
    implicit none
    integer loksvf,iws(*)
-----
    subroutine biblioread(bibhead,iws)
! read references from save file
    implicit none
    integer bibhead,iws(*)
-----
    subroutine readash(lok,iws)
! reading assessment records
    integer lok,iws(*)

```

8.9.5 Reinitiate workspaces by deleting exiting data

This is necessary before reading a new data from a database or unformatted save file.

```
subroutine new_gtp
!
! DELETES ALL DATA so a new TDB file can be read
!
! this is needed before reading a new unformatted file (or same file again)
! we must go through all records and delete and deallocate each
! separately. Very similar to gtpread
  implicit none
```

8.9.6 Delete data

This subroutine deletes all allocated data. Used when giving the NEW command. It does not deallocate data that has been allocated for step and map, that should be done beforehand.

```
subroutine deallocate_gtp(intvar,dblvar)
! deallocate the data structure
  implicit none
  integer allocatestatus
  integer intvar(*)
  double precision dblvar(*)
```

8.9.7 Delete a phase

```
subroutine delphase(lokph)
! save data for phase at location lokph (except data in the equilibrium record)
! For phases with disordered set of parameters we must access the number of
! sublattices via firsteq
  implicit none
  integer lokph
```

8.9.8 Yet another utility routine

Some utilities used by the routines reading database files.

```
logical function iskeyword(text,keyword,nextc)
! compare a text with a given keyword. Abbreviations allowed
! but the keyword and abbreviation must be surrounded by spaces
! nextc set to space character in text after the (abbreviated) keyword
  implicit none
```



```

character text*(*),keyword*(*),key*64
integer nextc
-----
integer function istdbkeyword(text,nextc)
! compare a text with a given keyword. Abbreviations allowed (not within _)
! but the keyword and abbreviation must be surrounded by spaces
! nextc set to space character in text after the (abbreviated) keyword
  implicit none
  character text*(*),keyword*(*),key*64
  integer nextc
  -----
integer function ispdbkeyword(text,nextc)
! compare a text with a given keyword. Abbreviations allowed (not within _)
! but the keyword and abbreviation must be surrounded by spaces
! nextc set to space character in text after the (abbreviated) keyword
  implicit none
  character text*(*),keyword*(*),key*64
  integer nextc
  -----
subroutine replacetab(line,nl)
! replaces TAB by space in line
  implicit none
  character line*(*),key*64
  integer nl

```

8.9.9 Read a TDB file

This subroutine can read a TDB file that is not too fancily edited manually. The best is to read a file written from Thermo-Calc. Some TYPE_DEFINITIONS are not handled. All TYPE_DEFINITIONS should be in the beginning of the TDB file, especially DISORDERD.PART as this has been implemented differently in OC.

```

subroutine readtdb(filename,nel,selel)
! reading data from a TDB file with selection of elements
!-----
! Not all TYPE_DEFS implemented
!-----
  implicit none
  integer nel
  character filename*(*),selel(*)*2
  -----
subroutine readtdbsilent
-----
subroutine any_disordered_part(lin,ndisph,disph)
! reading data from a PDB file with selection of elements

```

```

!-----
! Not all TYPE_DEFS implemented
!-----
      implicit none
      integer lin,ndisph
      character disph(*)*(*)

```

8.9.10 Reading a database in PDB format

A new format for thermodynamic databases is under development.

```

subroutine readpdb(filename,nel,selel,options)
! reading data from a PDB file with selection of elements
      implicit none
      integer nel
      character filename*(*),selel(*)*2,options*(*)

```

8.9.11 Writing a database in PDB format

NOT IMPLEMENTED. A new format for thermodynamic databases is under development.

```

subroutine write_pdbformat(unit)
! write a PDB database
      implicit none
      integer unit

```

8.9.12 Check a TDB file exists and extract elements

This is used to check a user typed a correct TDB file name and extracts the elements so the user can select which he wants.

```

subroutine checkdb(filename,ext,nel,selel)
! checking a TDB/PDB file exists and return the elements
      implicit none
      integer nel
      character filename*(*),ext*4,selel(*)*2
-----
subroutine checkdb2(filename,ext,nel,selel)
! checking a TDB/PDB file exists and return the elements
! It also writes 15 lines from any "DATABASE_INFO" in the file
      implicit none
      integer nel
      character filename*(*),ext*4,selel(*)*2

```

8.9.13 Save data in SOLGASMIX dat format

This routine will attempt to save the current data in OC on a file in the DAT format used by SOLGASMIX. There are also some subroutines handling TP functions involved and they are described in section 8.19.17.

```
subroutine save_datformat(filename,version,kod,ceq)
! writes a SOLGASMIX DAT format file. not (ever?) finished
  implicit none
  integer kod
  character filename*(*),version*(*)
  type(gtp_equilibrium_data), pointer :: ceq
-----
subroutine incunique(text)
  character text*(*)
-----
subroutine expand_wildcards(intconst,nconst,wildloop,iset,lokph)
! Expand a wildcard constituent with all constituents it replaces
! There can be several wildcards
! intconst is the original set of constituents including the wildcards (-99)
! nconst is the number of constituents
! wildloop is set to the number of times the interaction is repeated
! iset is a matrix with the expanded constituents
! phrecord is the phase record where one can find the phase structure
  implicit none
  integer intconst(*)
  integer, allocatable, dimension(:,:) :: iset
  integer nconst,wildloop,lokph
-----
subroutine intsort(intc,nint,intx)
! This is just another stupid sorting subroutine
! intc is not changed
  implicit none
  integer intc(*),intx(*),nint
-----
subroutine lower_case_species_name(constext,ip,isp)
! writes a species name using lower case for second letter of element
  implicit none
  character constext*(*)
  integer ip,isp
```

8.10 State variable stuff

State variables are important for the setting and extracting results of a calculation. State variables are treated very similarly to Thermo-Calc using symbols like T , P , N , $x(< \text{component} >)$ etc.

The internal syntax of state variables is rather complicated, perhaps it should be revised and defined as a structure? If there are errors or one wants to make modifications it is not easy.

Things like Curie temperature, Debye temperature, mobilities etc are tread almost like “state variables” although one cannot use them in conditions. Adding more things like elastic constants will be a bit complicated.

The subroutines for manipulations is also a bit complicated and could do with a clean up and renaming.

8.10.1 Get state variable value given its symbol

By providing a state variable as a character variable like T or $x(\text{liquid}, \text{cr})$ this routine returns its current value. Wildcards, “*”, are not allowed, see 8.10.2.

A variant checks if a specified phase is stable.

```

subroutine get_stable_state_var_value(statevar,value,encoded,ceq)
! called with a state variable character
! If the state variable includes a phase it checks if the phase is stable ...
  implicit none
  TYPE(gtp_equilibrium_data), pointer :: ceq
  character statevar*(*),encoded*(*)
  double precision value
-----
subroutine get_state_var_value(statevar,value,encoded,ceq)
! called with a state variable character
  implicit none
  TYPE(gtp_equilibrium_data), pointer :: ceq
  character statevar*(*),encoded*(*)
  double precision value

```

8.10.2 Get many state variable values

This routine can be called with wildcard, “*”, as argument in state variables like $NP(*)$, $x(*, CR)$ etc. It is fragile and currently only available when defining plot axis and some output.

```

subroutine get_many_svar(statevar,values,mjj,kjj,encoded,ceq)
! called with a state variable name with wildcards allowed like NP(*), X(*,CR)
! mjj is dimension of values, kjj is number of values returned
! encoded used to specify if phase data in phasetuple order ('Z')
! >>>> BIG question: How to do with phases that are note stable?
! If I ask for w(*,Cr) I only want the fraction in stable phases
! but when this is used for GNUMPLOT the values are written in a matix
! and the same column in that phase must be the same phase ...
! so I have to have the same number of phases from each equilibria.

```



```

! Typically: T, x(fe), x(fcc,fe), np(fcc), y(fcc,c#2), ac(h2,bcc), ac(fe)
! NOTE! model properties like TC(FCC),MQ&FE(FCC,CR) must be detected
! NOTE: added storing information in a gtp_state_variable record svrec !!
!
! this routine became as messy as I tried to avoid
! but I leave it to someone else to clean it up ...
!
! state variable and indices
! Symbol  no    index1 index2 index3 index4
! T       1     -
! P       2     -
! MU      3     component or phase,constituent
! AC      4     component or phase,constituent
! LNAC    5     component or phase,constituent
!
!                                     index (in svid array)
! U       10    (phase#set)           6     Internal energy (J)
! UM      11    "                     6     per mole components
! UW      12    "                     6     per kg
! UV      13    "                     6     per m3
! UF      14    "                     6     per formula unit
! S       2x    "                     7     entropy
! V       3x    "                     8     volume
! H       4x    "                     9     enthalpy
! A       5x    "                    10     Helmholtz energy
! G       6x    "                    11     Gibbs energy
! NP      7x    "                    12     moles of phase
! BP      8x    "                    13     mass of moles
! DG      9x    "                    15     Driving force
! Q      10x    "                    14     Internal stability
! N      11x    (component/phase#set,component) 16 moles of components
! X      111    "                    17     mole fraction of components
! B      12x    "                    18     mass of components
! W      122    "                    19     mass fraction of components
! Y       13    phase#set,constituent#subl 20     constituent fraction
!----- model variables <<<< these now treated differently
! TC      -     phase#set             -     Magnetic ordering T
! BMAG    -     phase#set             -     Aver. Bohr magneton number
! MQ&     -     element, phase#set    -     Mobility
! THET    -     phase#set             -     Debye temperature
!
implicit none
integer, parameter :: noos=20
character*4, dimension(noos), parameter :: svid = &
  ['T  ', 'P  ', 'MU ', 'AC  ', 'LNAC', 'U   ', 'S   ', 'V   ', '&
  'H   ', 'A   ', 'G   ', 'NP  ', 'BP  ', 'DG  ', 'Q   ', 'N   ', '&
  'X   ', 'B   ', 'W   ', 'Y   ' ]

```

```

!           1           2           3           4           4           6           7           8
character statevar*(*)
integer istv,iref,iunit
integer, dimension(4) :: indices
type(gtp_equilibrium_data), pointer :: ceq
! I shall try to use this record type instead of separate arguments: !!
!   type(gtp_state_variable), pointer :: svrec
!   type(gtp_state_variable), pointer :: svr

```

8.10.4 Calculate molar and mass properties for a phase

This subroutine calculates mole and mass fractions of all components for a phase (mole fractions of components not dissolved is zero). It also returns the total number of moles of components and the mass. In amount the number of moles per formula unit is returned (same as qq(1) in get_phase_data and set_constitution).

```

subroutine calc_phase_molmass(iph,ics,xmol,wmass,totmol,totmass,amount,ceq)
! calculates mole fractions and mass fractions for a phase#set
! xmol and wmass are fractions of components in mol or mass
! totmol is total number of moles and totmass total mass of components.
! amount is number of moles of components per formula unit.
implicit none
TYPE(gtp_equilibrium_data), pointer :: ceq
integer iph,ics
double precision, dimension(*) :: xmol,wmass
double precision amount,totmol,totmass

```

8.10.5 Calculate molar amounts for a phase

Specially used for grid minimization.

```

subroutine calc_phase_mol(iph,xmol,ceq)
! calculates mole fractions for phase iph, compset 1 in equilibrium ceq
! used for grid generation and some other things
! returns current constitution in xmol equal to mole fractions of components
implicit none
integer iph
double precision xmol(*)
TYPE(gtp_equilibrium_data),pointer :: ceq

```

8.10.6 Sum molar and mass properties for all phases

Sums the mole and mass fractions for all components and also total number of moles and mass over all stable phases using 8.10.4.

```

subroutine calc_molmass(xmol,wmass,totmol,totmass,ceq)
! summing up N and B for each component over all phases with positive amount
! Check that totmol and totmass are correct ....
  implicit none
  double precision, dimension(*) :: xmol,wmass
  double precision totmol,totmass
  TYPE(gtp_equilibrium_data), pointer :: ceq

```

8.10.7 Sum all normalizing property values

Used to calculate normalizing properties like V, N and B but also G and S for the whole system.
Used when calculating state variable values.

```

subroutine sumprops(props,ceq)
! summing up G, S, V, N and B for all phases with positive amount
! Check if this is correct
  implicit none
  TYPE(gtp_equilibrium_data), pointer :: ceq
  double precision props(5)

```

8.10.8 Encode state variable

This converts the internal coding of a state variable into a character variable text starting at position ip. The value of ip is updated inside.

The subroutine also handles property symbols used in the parameters, see 2.3.3, to make it possible to list the symbol of such a property after an equilibrium calculation. See 8.10.3

The new version of this subroutine uses state variable records, the previous one using individual arguments should not be used as it will eventually disappear.

```

subroutine encode_state_variable(text,ip,svr,ceq)
! writes a state variable in text form position ip. ip is updated
  character text*(*)
  integer ip
  type(gtp_state_variable), pointer :: svr
  type(gtp_equilibrium_data), pointer :: ceq
-----
subroutine encode_state_variable3(text,ip,istv,indices,iunit,iref,ceq)
! writes a state variable in text form position ip. ip is updated
! the internal coding provides in istv, indices, iunit and iref
! ceq is needed as components can be different in different equilibria ??
! >>>> unfinished as iunit and iref not really cared for
  implicit none
  integer, parameter :: noos=20

```



```

character*4, dimension(noos), parameter :: svid = &
  ['T  ','P  ','MU  ','AC  ','LNAC','U  ','S  ','V  ',&
   'H  ','A  ','G  ','NP  ','BP  ','DG  ','Q  ','N  ',&
   'X  ','B  ','W  ','Y  ' ]
character*(*) text
integer, dimension(4) :: indices
integer istv,ip,iunit,iref
type(gtp_equilibrium_data), pointer :: ceq

```

8.10.9 Encode a state variable record

This is provided to convert a single state variable record to a text.

```

subroutine encode_state_variable_record(text,ip,svr,ceq)
! writes a state variable in text form position ip. ip is updated
! the svr record provide istv, indices, iunit and iref
! ceq is needed as compopnents can be different in different equilibria ??
! >>>> unfinished as iunit and iref not really cared for
implicit none
integer, parameter :: noos=20
character*4, dimension(noos), parameter :: svid = &
  ['T  ','P  ','MU  ','AC  ','LNAC','U  ','S  ','V  ',&
   'H  ','A  ','G  ','NP  ','BP  ','DG  ','Q  ','N  ',&
   'X  ','B  ','W  ','Y  ' ]
character*(*) text
type(gtp_state_variable) :: svr
type(gtp_equilibrium_data), pointer :: ceq

```

8.10.10 Calculate state variable value

This is the subroutine that actually calculates the value of a state variable. The state variable is identified using the internal coding.

The subroutine also handle properties used in the parameters, see 2.3.3, to make it possible to obtain the value of such a property after an equilibrium calculation. The values of istv etc must be as returned from decode_state_variable, see 8.10.3.

```

subroutine state_variable_val(svr,value,ceq)
! calculate the value of a state variable in equilibrium record ceq
! It transforms svr data to old format and calls state_variable_val3
type(gtp_state_variable), pointer :: svr
double precision value
TYPE(gtp_equilibrium_data), pointer :: ceq
-----
subroutine state_variable_val3(istv,indices,iref,iunit,value,ceq)

```

```

! calculate the value of a state variable in equilibrium record ceq
! istv is state variable type (integer)
! indices are possible specifiers
! iref indicates use of possible reference state, 0 current, -1 SER
! iunit is unit, (K, oC, J, cal etc). For % it is 100
! value is the calculated values. for state variables with wildcards use
! get_many_svar
  implicit none
  integer, dimension(4) :: indices
  TYPE(gtp_equilibrium_data), pointer :: ceq
  integer istv,iref,iunit
  double precision value

```

8.10.11 Calculate eigenvalues of a phase stability matrix

This subroutine calculates the eigenvalues of the phase stability matrix, i.e. the matrix of all second derivatives with respect to the constitution. It returns the value that is lowest, i.e. negative or closest to zero. If it is negative the phase composition is inside a spinodal. It uses a LINPAC routine to calculate the eigenvalues.

```

subroutine calc_qf(lokcs,value,ceq)
! calculates eigenvalues of the second derivative matrix, stability function
! lokcs is index of phase_varres
! value calculated value returned
! ceq is current equilibrium
! For ionic liquid and charged crystalline phases one should
! calculate eigenvectors to find neutral directions.
  implicit none
  TYPE(gtp_equilibrium_data), pointer :: ceq
  integer :: lokcs
  double precision value

```

8.10.12 The value of the user defined reference state

The value of many state variables depend on the selected reference state. By default that is the value at 298.15 K and 1 bar for the stable phase of the pure elements, this is called SER. If the user defines another reference state this subroutine calculates that value. The reference state is set calling the routine in section 8.5.2.

```

subroutine calculate_reference_state(kstv,iph,ics,aref,ceq)
! Calculate the user defined reference state for extensive properties
! kstv is the type of property: 1 U, 2 S, 3 V, 4 H, 5 A, 6 G
! It can be phase specific (iph.ne.0) or global (iph=0)
! IMPORTANT

```

```

! For integral quantities (like calculated here) the reference state
! is ignored unless all components have the same phase as reference (like Hmix)
  implicit none
  integer kstv,iph,ics
  double precision aref
  type(gtp_equilibrium_data), pointer :: ceq
-----
  subroutine calculate_reference_state_old(kstv,iph,ics,aref,ceq)
! Calculate the user defined reference state for extensive properties
! kstv is the type of property: 1 U, 2 S, 3 V, 4 H, 5 A, 6 G
! It can be phase specific (iph.ne.0) or global (iph=0)
! IMPORTANT
! For integral quantities (like calculated here) the reference state
! is ignored unless all components have the same phase as reference (like Hmix)
  implicit none
  integer kstv,iph,ics
  double precision aref
  type(gtp_equilibrium_data), pointer :: ceq

```

8.11 State variable functions

This section is separated from the state variables itself to make it a little simpler. State variable function can contain any combination of state variables using normal operators like +, -, *, / but also EXP, LN, LOG10, ERF etc. The PUTFUN subroutine in the METLIB package is used. No derivatives can be calculated. A state variable function can refer to another state variable function. Thus when the value of a state variable function is requested all are calculated with some exceptions: not dot derivatives and not those amended as:

- The function is amended to be evaluated only when specified explicitly. Such a symbol can be used as condition for example in STEP EVALUATE (not yet implemented).
- The function is amended to be evaluated only in a specific equilibrium record. This is useful if one has experiments of heat differences relative to an equilibrium at a certain T . The enthalpy of this equilibrium will be evaluated only at this equilibrium.

An extension planned but not yet implemented is to allow formal arguments when defining a state variable function, for example $CP(@P)=HM(@P).T$ where the formal argument @P means a phase. @S would stand for a species and @C for a component. When calling the function an actual argument must be supplied.

Some state variable functions using “dot derivatives” to calculate for example the heat capacity are calculated inside the minimizer routine as they need results for the last equilibrium.

8.11.1 Enter a state variable function

The first subroutine enters a state variable function and the other stores it in the SVFLISTA array. The old version will eventually disappear.

```
subroutine enter_svfun(cline,last,ceq)
! enter a state variable function
  implicit none
  integer last
  character cline*(*)
  type(gtp_equilibrium_data), pointer :: ceq
-----
subroutine set_putfun_constant(svfix,value)
! changes the value of a putfun constant
! svfix is index, value is new value
  implicit none
  integer svfix
  double precision value
-----
subroutine store_putfun(name,lrot,nsymb,iarr)
! enter an expression of state variables with name name with address lrot
! nsymb is number of formal arguments
! iarr identifies these
  implicit none
  character name*(*)
  type(putfun_node), pointer :: lrot
  integer nsymb
  integer iarr(10,*)
-----
subroutine store_putfun_old(name,lrot,nsymb,&
  istv,indstv,iref,iunit,idot)
! enter an expression of state variables
! name: character, name of state variable function
! lrot: pointer, to a putfun_node that is the root of the stored expression
! nsymb: integer, number of formal arguments
! istv: integer array, formal argument state variables typ
! indstv: 2D integer array, indices for the formal state variables
! iref: integer array, reference for the formal state variables
! iunit: integer array, unit of the formal state variables
  implicit none
  type(putfun_node), pointer :: lrot
  integer nsymb
  integer, dimension(*) :: istv,iref,iunit,idot
  integer, dimension(4,*) :: indstv
  character name*(*)
```

8.11.2 List a state variable function

These two subroutines can find a state variable function and list its name in a character respectively.

```
subroutine find_svfun(name,lrot)
! finds a state variable function called name (no abbreviations)
! ceq not needed!??
  implicit none
  character name*(*)
  integer lrot
!   type(gtp_equilibrium_data), pointer :: ceq
-----
subroutine find_symbol_with_equilno(lrot,eqno)
! finds a state variable function with equilibrium index
  implicit none
  integer lrot,eqno
-----
subroutine list_svfun(text,ipos,lrot,ceq)
! list a state variable function
  implicit none
  character text*(*)
  integer ipos,lrot
  type(gtp_equilibrium_data), pointer :: ceq
```

8.11.3 Utility subroutine for state variable functions

Utility to store state variable identification for a function.

```
subroutine make_stvrec(svr,iarr)
! stores appropriate values from a formal argument list to a state variable
! function in a state variable record
  implicit none
  type(gtp_state_variable), pointer :: svr
  integer iarr(10)
```

8.11.4 List all state variable functions

Lists all state variables functions on device kou.

```
subroutine list_all_svfun(kou,ceq)
! list all state variable funtions
  implicit none
  integer kou
  type(gtp_equilibrium_data), pointer :: ceq
```

8.11.5 Some depreciated routines

The double precision function `evaluate_svfun_old(lrot,actual_arg,mode,ceq)` is now in the minimizer package but still used somewhere.

```
subroutine evaluate_all_svfun_old(kou,ceq)
! THIS SUBROUTINE MOVED TO MINIMIZER but kept for initiallizing
! cannot be used for state variable functions that are derivatives ...
! evaluate and list values of all functions but it is still used somewhere
  implicit none
  integer kou
  TYPE(gtp_equilibrium_data), pointer :: ceq
-----
double precision function evaluate_svfun_old(lrot,actual_arg,mode,ceq)
! THIS SUBROUTINE MOVED TO MINIMIZER
! but needed in some cases in this module ... ???
! envaluate all funtions as they may depend on each other
! actual_arg are names of phases, components or species as @Pi, @Ci and @Si
! needed in some deferred formal parameters (NOT IMPLEMENTED YET)
! if mode=1 always evaluate
  implicit none
  integer lrot,mode
  character actual_arg(*)*(*)
  TYPE(gtp_equilibrium_data), pointer :: ceq
```

8.12 Status for things

There are many status bits in various records. These subroutines and function set, clear or test these.

Setting means to set the bit to 1 and clearing means to set the bit to 0. A problem is that I always mix up if BTEST return TRUE or FALSE if the bit is set.

The subroutines and functions in this section should be simplified.

8.12.1 Set status for elements

An element can be entered or suspended.

```
subroutine change_element_status(elname,nystat,ceq)
! change the status of an element, can affect species and phase status
! nystat:0=entered, 1=suspended, -1 special (exclude from sum of mole fraction)
!
! suspending elements for each equilibrium separately not yet implemented
!
```

```

implicit none
character elname*(*)
integer nystat
TYPE(gtp_equilibrium_data), pointer :: ceq

```

8.12.2 Test status for element

An element can be entered or suspended.

```

logical function testelstat(iel,status)
! return value of element status bit
implicit none
integer iel,status

```

8.12.3 Set status for species

A species can be entered or suspended.

```

subroutine change_species_status(spname,nystat,ceq)
! change the status of a species, can affect phase status
! nystat:0=entered, 1=suspended
implicit none
integer nystat
character spname*(*)
TYPE(gtp_equilibrium_data), pointer :: ceq

```

8.12.4 Test status for species

A species can be entered or suspended.

```

logical function testspstat(isp,status)
! return value of species status bit
implicit none
integer isp,status

```

8.12.5 Get and test status for phase

Get and test phase status are almost identical but the get routine returns some additional information. Note that there are also phase bits that determine the model used.

```

integer function get_phase_status(ip,ics,text,ip,val,ceq)
! return phase status as text and amount formula units in val

```

```

! for entered and fix phases also phase amounts.
! OLD Function value: 1=entered, 2=fix, 3=dormant, 4=suspended, 5=hidden
  implicit none
  character text*(*)
  integer iph,ics,ip
  TYPE(gtp_equilibrium_data), pointer :: ceq
  double precision val
-----
  integer function test_phase_status(iph,ics,val,ceq)
! Almost same as get_..., returns phase status as function value but no text
! old: 1=entered, 2=fix, 3=dormant, 4=suspended, 5=hidden
! nystat:-4 hidden, -3 suspended, -2 dormant, -1,0,1 entered, 2 fix
! this is different from in change_phase .... one has to make up one's mind
  implicit none
  TYPE(gtp_equilibrium_data), pointer :: ceq
  integer iph,ics
  double precision val

```

8.12.6 Set/unset/test phase model bit

This sets bits indicating if the phase is ideal, has fcc ordering etc. It does not test if ENTERED/FIXED/DORMANT or SUSPENDED.

```

  subroutine set_phase_status_bit(lokph,bit)
! set the status bit "bit" in status1, cannot be done outside this module
! as the phlista is private
! These bits do not depend on the composition set
  implicit none
  integer lokph,bit
-----
  subroutine clear_phase_status_bit(lokph,bit)
! clear the status bit "bit" in status1, cannot be done outside this module
! as the phlista is private
  implicit none
  integer lokph,bit
-----
  logical function test_phase_status_bit(iph,ibit)
! return TRUE is status bit ibit for phase iph, is set
! because phlista is private. Needed to test for gas, ideal etc,
! DOES NOT TEST STATUS like entered/fixed/dormant/suspended
  implicit none
  integer iph,ibit

```


8.12.7 Change status for phase

Several subroutines to change the status bits for one or more phases. Note NYSTAT values are different from this in GET and TEST above. Same values should be used.

```
subroutine change_many_phase_status(phnames,nystat,val,ceq)
! change the status of many phases.
! nystat:-4 hidden, -3 suspended, -2 dormant, -1,0,1 entered, 2 fix
! phnames is a list of phase names or *S (all suspended) *D (all dormant) or
! *E (all entered (stable, unknown, unstable), *U all unstable
! If just * then change_phase_status is called directly
! It calls change_phase_status for each phase
  implicit none
  character phnames*(*)
  integer nystat
  double precision val
  TYPE(gtp_equilibrium_data), pointer :: ceq
-----
subroutine change_phtup_status(phtupx,nystat,val,ceq)
! change the status of a phase tuple. Also used when setting phase fix etc.
! nystat:-4 hidden, -3 suspended, -2 dormant, -1,0,1 entered, 2 fix
! qph can be -1 meaning all or a specifix phase index. ics compset
!
  implicit none
  integer phtupx,nystat
  double precision val
  TYPE(gtp_equilibrium_data), pointer :: ceq
-----
subroutine change_phase_status(qph,ics,nystat,val,ceq)
! change the status of a phase. Also used when setting phase fix etc.
! nystat:-4 hidden, -3 suspended, -2 dormant, -1,0,1 entered, 2 fix
! qph can be -1 meaning all or a specifix phase index. ics compset
!
  implicit none
  integer qph,ics,nystat
  double precision val
  TYPE(gtp_equilibrium_data), pointer :: ceq
-----
subroutine mark_stable_phase(iph,ics,ceq)
! change the status of a phase. Does not change fix status
! called from meq_sameset to indicate stable phases (nystat=1)
! nystat:-4 hidden, -3 suspended, -2 dormant, -1,0,1 entered, 2 fix
!
  implicit none
  integer iph,ics
  TYPE(gtp_equilibrium_data), pointer :: ceq
```

8.12.8 Set unit for energy etc.

This is not yet implemented.

```
subroutine set_unit(property,unit)
! set the unit for a property, like K, F or C for temperature
! >>>> unfinished
  implicit none
  character*(*) property,unit
```

8.12.9 Save results for a phase

This is not yet implemented.

```
subroutine save_results(lut,iph,ics,long)
! write calculated results for a phase for later use in POST
  implicit none
  integer lut,iph,ics,long
```

8.12.10 Set reference state for constituent

This is not yet implemented.

```
subroutine set_constituent_reference_state(iph,icon,asum)
! determine the end member to calculate as reference state for this constituent
! Used when giving a chemical potential for a constituent like MU(GAS,H2O)
  implicit none
  integer iph,icon
  double precision asum
```

8.12.11 Calculate conversion matrix for new components

```
subroutine elements2components1(nspel,dum,ncmp,cmpstoi,ceq)
! converts a stoichiometry array for a species from elements to components
! This subroutine, is it used to get activity for a constituent in gtp3F
! dum is is no longer used
  implicit none
  integer nspel,ncmp
! cmpstoi is stoichiometry as element, changed to be as components
  double precision cmpstoi(*),dum(*)
  double precision, allocatable :: stoi(:)
  type(gtp_equilibrium_data), pointer :: ceq
```

8.13 Internal stuff

The subroutines in this section are internal and not to be used by calls from outside GTP.

8.13.1 Indicate the type of an experiment

An experiment can specify an equal value, =, a larger than limit, > or a lesser than limit, <. This return the type.

```
subroutine termterm(string,ich,kpos,lpos,value)
! search for first occurrence of + - = > or <
! if + or - then also extract possible value after sign
! value is coefficient for NEXT term (if any)
! kpos is last character in THIS state variable, lpos where NEXT may start
  implicit none
  character string*(*)
  integer kpos,ich,lpos
  double precision value
```

8.13.2 Alphabetical ordering

The elements, species and phases are stored in the arrays ELLISTA, SPLISTA and PHLISTA in the order they are entered. They are sorted in alphabetical order in the array ELEMENTS, SPECIES and PHASES. These routines are used to maintain the alphabetical order.

```
subroutine alphaelorder
! arrange new element in alphabetical order
! also make alphaindex give alphabetical order
  implicit none
-----
subroutine alphasporder
! arrange new species in alphabetical order
! also make alphaindex give alphabetical order
  implicit none
-----
subroutine alphaphorder(tuple)
! arrange last added phase in alphabetical order
! also make alphaindex give alphabetical order
! phletter G and L and I have priority
! tuple is returned as position in phase tuple
  implicit none
  integer tuple
```

8.13.3 Check alphaindex

Just for testing.

```
subroutine check_alphaindex
! just for debugging, check that ellist(i)%alphaindex etc is correct
implicit none
```

8.13.4 Creates a list of constituents of a phase

Not much to discuss.

```
subroutine create_constitlist(constitlist,nc,klist)
! creates a constituent list ...
implicit none
integer, dimension(*) :: klist
integer, dimension(:), allocatable :: constitlist
integer nc
```

8.13.5 Creates a new parrecord for a phase

Not much to discuss either.

```
subroutine create_parrecords(lokph,lokcs,ns1,nc,nprop,iva,ceq)
! fractions and results arrays for a phase for parallel calculations
! location is returned in lokcs
! ns1 is sublattices, nc number of constituents, nprop max number if propt,
! iva is an array which is set as constituent status word (to indicate VA)
! ceq is always firsteq ???
!
! BEWARE not adopted for threads
!
! >>> changed all firsteq below to ceq????
!
implicit none
TYPE(gtp_equilibrium_data), pointer :: ceq
integer, dimension(*) :: iva
integer lokph,lokcs, ns1, nc, nprop
```

8.13.6 Create interaction record

Finds the correct place to add an interaction parameter. It should always be linked from the first possible endmember (with the alphabetically first constituents) and then in alphabetical order of the interaction elements.

```

subroutine create_interaction(intrec,mint,lint,intperm,intlinks)
! creates a parameter interaction record
! with permutations if intperm(1)>0
  implicit none
  type(gtp_interaction), pointer :: intrec
  integer, dimension(2,*) :: lint,intlinks
  integer, dimension(*) :: intperm
  integer mint

```

8.13.7 Create endmember record

Creates a record for an endmember. Sometimes endmembers have no property records if there are interaction parameters must be linked from this endmember.

```

subroutine create_endmember(lokph,newem,noperm,nsl,endm,elinks)
! create endmember record with nsl sublattices with endm as constituents
! noperm is number of permutations
! endm is the basic endmember (if there are permutations)
! elinks are the links to constituents for all permutations
  implicit none
  integer endm(*)
  integer lokph,noperm,nsl
  type(gtp_endmember), pointer :: newem
  integer, dimension(nsl,noperm) :: elinks

```

8.13.8 Create and extend a property record

All parameter values for an endmember or interaction record are stored in property records. An endmember or interaction parameter may have several property records linked in a list.

An interaction record can have a degree and each degree a function. When entering a function for a higher degree than in the current property record it must be extended.

```

subroutine create_proprec(proprec,proptype,degree,lfun,refx)
! reserves a property record from free list and insert data
  implicit none
  TYPE(gtp_property), pointer :: proprec
  integer proptype,degree,lfun
  character refx*(*)
  -----
{\small
\begin{verbatim}
  subroutine extend_proprec(current,degree,lfun)
! extends a property record and insert new data
  implicit none

```

```

integer degree,lfun
type(gtp_property), pointer :: current

```

8.13.9 Add a fraction set record

```

subroutine add_fraction_set(iph,id,ndl,totdis)
! add a new set of fractions to a phase, usually to describe a disordered state
! like the "partitioning" in old TC
!
! BEWARE this is only done for firsteq, illegal when having more equilibria
!
! id is a letter used as suffix to identify the parameters of this set
! ndl is the last original sublattice included in the (first) disordered set
! ndl can be 1 meaning sublattice 2..nsl are disordered, or nsl meaning all are
!   disordered
! totdis=0 if phase never disorder totally (like sigma)
!
! For a phase like (Al,Fe,Ni)3(Al,Fe,Ni)1(C,Va)4 to add (Al,Fe,Ni)4(C,Va)4
! icon=1 2 3 1 2 3 4 5 with ndl=2
! For a phase like (Fe,Ni)10(Cr,Mo)4(Cr,Fe,Mo,Ni)16 then
! icon=2 4 1 3 1 2 3 4 with ndl=3
! This subroutine will create the necessary data to calculate the
! disordered fraction set from the site fractions.
!
! IMPORTANT (done): for each composition set this must be repeated
! if new composition sets are created it must be repeated for these
!
! IMPORTANT (not done): order the constituents alphabetically in each disorderd
! sublattice otherwise it will not be possible to enter parameters correctly
!
  implicit none
  integer iph,ndl,totdis
  character id*1

```

8.13.10 Copy record for fraction sets

```

subroutine copy_fracset_record(lokcs,disrec,ceq)
! attempt to create a new disordered record ??? this can probably be done
! with just one statement .. but as it works I am not changing right now
  implicit none
  TYPE(gtp_equilibrium_data), pointer :: ceq
  TYPE(gtp_fraction_set) :: disrec
  integer lokcs

```

8.13.11 Implicit suspend and restore

If an element is suspended some species may have to be suspended too and if a species is suspended some phases may have to be suspended. This routine does that.

```
subroutine suspend_species_implicitly(ceq)
! loop through all entered species and suspend those with an element suspended
  implicit none
  TYPE(gtp_equilibrium_data), pointer :: ceq
-----
subroutine suspend_phases_implicitly(ceq)
! loop through all entered phases and suspend constituents and
! SUSPEND phases with all constituents in a sublattice suspended
!   dimension lokcs(9)
  implicit none
  TYPE(gtp_equilibrium_data), pointer :: ceq
-----
subroutine restore_species_implicitly_suspended
! loop through all implicitly suspended species and restore those with
! all elements entered
  implicit none
-----
subroutine restore_phases_implicitly_suspended
! loop through all implicitly suspended phases and restore those with
! at least one constituent entered in each sublattice
  implicit none
```

8.13.12 Add to reference phase

There is a reference phase that should have parameters for each element in its stable state at all temperatures and 1 bar. For each element entered this subroutine adds it to the reference phase. This phase can never be used in calculations. It represents different phases for each element, gas for H, bcc for Cr etc.

```
subroutine add_to_reference_phase(loksp)
! add this element to the reference phase
! loksp: species index of new element
  implicit none
  integer loksp
```

8.14 Additions

Creating, handling and calculations of additions to the Gibbs energy. This is a section that will probably be extended with several new subroutines for different kinds of additions. See also sections 1.4, 3.3.1 and 2.3.3,

Note that the Equi-entropy criterion (EET) is checked in the minimizer as it relates to two different phases (a solid and a liquid). It is an addition as it does not depend on property of a single phase.

```

subroutine addition_selector(addrrec,moded,phres,lokph,mc,ceq)
! called when finding an addition record while calculating G for a phase
! addrrec is addition record
! moded is 0, 1 or 2 if no, first or 2nd order derivatives should be calculated
! phres is ?
! lokph is phase location
! mc is number of constitution fractions
! ceq is current equilibrium record
  implicit none
  type(gtp_phase_add), pointer :: addrrec
  integer moded,lokph,mc
  TYPE(gtp_phase_varres), pointer :: phres
  type(gtp_equilibrium_data), pointer :: ceq

```

8.14.1 Generic subroutine to add an addition

Not very well structured here.

```

subroutine add_addrecord(lokph,extra,addtyp)
! generic subroutine to add an addition typ addtyp (including Inden)
  implicit none
  integer lokph,addtyp
  character extra*(*)

```

8.14.2 Utility routines for addition

Searches for the composition dependent properties for a specific addition.

The subroutine “setpermolebit” indicates that the property should be calculated per mole of atoms, not per formula unit as default.

```

subroutine need_propertyid(id,typty)
! get the index of the property needed
  implicit none
  integer typty
  character*4 id
-----
subroutine setpermolebit(lokph,addtype)
! set bit in addition record that addition is per mole
! lokph is phase record

```



```

! addtype is the addition record type
  implicit none
  integer lokph,addtype

```

8.14.3 Enter and calculate Inden magnetic model

The first two subroutines create the ferromagnetic addition due to Inden model and store all necessary data inside this. The last subroutine is called when calculating the Gibbs energy for a phase if there is a magnetic addition linked to the phase. It must calculate the contribution to G and all first and second derivatives of G. The model parameter identifiers TC and BMAG defined in Table 1 are used.

In the call the pointer to the phase_varres record is provided where current values of G and derivatives can be found. Values of the ferromagnetic temperature and its derivatives with respect to constitution is also stored there. The chain rule for derivatives must be applied.

```

subroutine create_magrec_inden(addrrec,aff)
! enters the magnetic model
  implicit none
  type(gtp_phase_add), pointer :: addrrec
  integer aff
-----
subroutine calc_magnetic_inden(moded,phres,lokadd,lokph,mc,ceq)
! calculates Indens magnetic contribution
! NOTE: values for function not saved, should be done to save time.
! Gmagn = RT*f(T/Tc)*ln(beta+1)
! moded: integer, 0=only G, S, Cp; 1=G and dG/dy; 2=Gm dG/dy and d2G/dy2
! phres: pointer, to phase\_varres record
! lokadd: pointer, to addition record
! lokph: integer, phase record
! mc: integer, number of constituents
! ceq: pointer, to gtp\_equilibrium\_data
  implicit none
  integer moded,lokph,mc
  TYPE(gtp_phase_varres) :: phres
  TYPE(gtp_phase_add), pointer :: lokadd
  TYPE(gtp\_equilibrium\_data), pointer :: ceq

```

8.14.4 Create new magnetic model

Wei Xiagong has proposed a more advanced magnetic model. It is not yet fully implemented. It should use several model parameter identifies in Table 1

```

subroutine create_xiongmagnetic(addrrec,lokph,bcc)
! adds a Xiong type magnetic record, we must separate fcc and bcc by extra

```

```

! copied from Inden magnetic model
! The difference is that it uses TCA for Curie temperature and TNA for Neel
! and individual Bohr magneton numbers
  implicit none
  logical bcc
  integer lokph
  type(gtp_phase_add), pointer :: addrec
-----
  subroutine calc_xiongmagnetic(moded,phres,lokadd,lokph,mc,ceq)
! calculates Inden-Qing-Xiong magnetic contribution
!
! Gmagn = RT*f(T/Tc)*ln(beta+1)
! moded: integer, 0=only G, S, Cp; 1=G and dG/dy; 2=Gm dG/dy and d2G/dy2
! phres: pointer, to phase\_varres record
! lokadd: pointer, to addition record
! lokph: integer, phase record
! mc: integer, number of constituents
! ceq: pointer, to gtp\_equilibrium\_data
  implicit none
  integer moded,lokph,mc
! phres points to result record with gval etc for this phase
  TYPE(gtp_phase_varres) :: phres
  TYPE(gtp_phase_add), pointer :: lokadd
  TYPE(gtp\_equilibrium\_data), pointer :: ceq

```

8.14.5 Calculate and calculate volume model

A very simple volume model with two model parameters, V0 which is the volume at 298.15 K and 1 bar for the endmember, and VA which is the thermal expansion of the endmember defined in Table 1.

```

  subroutine create_volmod1(addrec)
! create addition record for the simple volume model
!
! currently only V = V0 * exp(VA(T))
! V0 is property (typty) 21, VA is 22 and reserved VB (Bulk modulus) 23
! but VB is not implemented yet
  implicit none
  type(gtp_phase_add), pointer :: addrec
-----
  subroutine calc_volmod1(moded,phres,lokadd,lokph,mc,ceq)
! calculate the simple volume model, CURRENTLY IGNORING COMPOSITION DEPENDENCE
!
! G = P*V0(x)*exp(VA(T,x))
! moded: integer, 0=only G, S, Cp; 1=G and dG/dy; 2=Gm dG/dy and d2G/dy2

```

```

! phres: pointer, to phase\_varres record
! lokadd: pointer, to addition record
! lokph: integer, phase record index
! mc: integer, number of constituents
! ceq: pointer, to gtp\_equilibrium\_data
  implicit none
  integer moded,lokph,mc
! phres points to result record with gval etc for this phase
  TYPE(gtp\_phase\_varres) :: phres
  TYPE(gtp\_phase\_add), pointer :: lokadd
  TYPE(gtp\_equilibrium\_data), pointer :: ceq

```

8.14.6 Calculate and calculate elastic contribution

This should create and calculate the elastic record. It is not implemented but there are several model parameters such as LPX, LPY, LPZ LPTH, EC11, EC12, EC44 previewed in Table 1.

```

subroutine create_elastic_model_a(newadd)
! addition record to calculate the elastic energy contribution
  implicit none
  type(gtp\_phase\_add), pointer :: newadd
-----
subroutine calc_elastica(moded,phres,addrec,lokph,mc,ceq)
! calculates elastic contribution and adds to G and derivatives
  implicit none
  integer moded,lokph,mc
  type(gtp\_phase\_varres), pointer :: phres
  type(gtp\_phase\_add), pointer :: addrec
  type(gtp\_equilibrium\_data), pointer :: ceq
-----
subroutine set_lattice_parameters(iph,ics,xxx,ceq)
! temporary way to set current lattice parameters for use with elastic model a
  implicit none
  integer iph,ics
  double precision, dimension(3,3) :: xxx
  type(gtp\_equilibrium\_data), pointer :: ceq

```

8.14.7 Heat capacity model for Einstein solids

This is now implemented. The parameter identifier THET in Table 1 is the logarithm of the Einstein T as that should be the reasonable property varying with the composition. See section 2.3.6.

```

subroutine create_einsteincp(newadd)
  implicit none

```

```

        type(gtp_phase_add), pointer :: newadd
-----
        subroutine calc_einsteincp(moded,phres,addrec,lokph,mc,ceq)
! Calculate the contibution due to Einste Cp model for low T
! moded 0, 1 or 2
! phres all results
! addrec pointer to addition record
! lokph phase record
! mc number of variable fractions
! ceq equilibrium record
!
!  $G = 1.5 \cdot R \cdot \text{THET} + 3 \cdot R \cdot T \cdot \ln(1 - \exp(-\text{THET}/T))$ 
! This is easier to handle inside the calc routine without TPFUN
!
        implicit none
        integer moded,lokph,mc
        type(gtp_phase_varres), pointer :: phres
        type(gtp_phase_add), pointer :: addrec
        type(gtp_equilibrium_data), pointer :: ceq

```

8.14.8 Schottky anomaly

This can describe a Schottky anomaly i.e. an increase of the heat capacity over a range in T followe by a decrease. It depends on the model parameter identifiers TSCH and CSCH, defined in Table 1.

```

        subroutine create_schottky_anomaly(newadd)
! Adding a Schottky anomaly to Cp
        implicit none
        type(gtp_phase_add), pointer :: newadd
-----
        subroutine calc_schottky_anomaly(moded,phres,addrec,lokph,mc,ceq)
! Calculate the contibution due to a Schottky anomaly
! moded 0, 1 or 2
! phres all results
! addrec pointer to addition record
! lokph phase record
! mc number of variable fractions
! ceq equilibrium record
!
!  $G = DCP2 \cdot T \cdot \ln(1 + \exp(-\text{THT2}/T))$ 
!  $dG/dT = DCP2 \cdot (\ln(1 + \exp(-\text{THT2}/T)) + (\text{THT}/T) \cdot (1 + \exp(+\text{THT2}/T))^{*-1})$ 
!  $d^2G/dT^2 = -DCP2 \cdot \text{THT2}^{**2} \cdot T^{*(-3)} \cdot \exp(\text{THT2}/T) \cdot (1 + \exp(+\text{THT2}/T))^{*(-2)}$ 
!
        implicit none

```

```

integer moded,lokph,mc
type(gtp_phase_varres), pointer :: phres
type(gtp_phase_add), pointer :: addrec
type(gtp_equilibrium_data), pointer :: ceq

```

8.14.9 Second Einstein

This can describe an smooth increase of the heat capacity to new fix value over a range in T . It depends on the model parameter identifiers THT2 and DCP2, defined in Table 1.

```

subroutine create_secondeinstein(newadd)
  implicit none
  type(gtp_phase_add), pointer :: newadd
  -----
  subroutine calc_secondeinstein(moded,phres,addrec,lokph,mc,ceq)
  ! Calculate the contibution due to Einste Cp model for low T
  ! moded 0, 1 or 2
  ! phres all results
  ! addrec pointer to addition record
  ! lokph phase record
  ! mc number of variable fractions
  ! ceq equilibrium record
  !
  ! G = 1.5*R*THET + 3*R*T*ln( 1 - exp( -THET/T ) )
  ! This is easier to handle inside the calc routine without TPFUN
  !
  implicit none
  integer moded,lokph,mc
  type(gtp_phase_varres), pointer :: phres
  type(gtp_phase_add), pointer :: addrec
  type(gtp_equilibrium_data), pointer :: ceq

```

8.14.10 Glass addition

Now it is implemented. The low T liquid is treated as an Einstein solid with an Einstein θ and an possible T dependent terms in the low T range. An additional model parameter, G2, defined in Table 1 will create an increase in the heat capacity before melting.

```

subroutine create_twostate_model1(addrec)
! newadd is location where pointer to new addition record should be stored
  implicit none
  type(gtp_phase_add), pointer :: addrec
  -----
  subroutine calc_twostate_model_john(moded,phres,addrec,lokph,mc,ceq)

```

```

! subroutine calc_twostate_model1(moded,phres,addrec,lokph,mc,ceq)
! This routine works OK but I am testing a modification
! moded is 0, 1 or 2 if no, first or 2nd order derivatives should be calculated
! addrec is addition record
! phres is phase_varres record
! lokph is phase location
! mc is number of constitution fractions
! ceq is current equilibrium record
  implicit none
  integer moded,lokph,mc
  TYPE(GTP_PHASE_ADD), pointer :: addrec
  TYPE(GTP_PHASE_VARRES), pointer :: phres
  TYPE(GTP_EQILIBRIUM_DATA), pointer :: ceq
-----
  subroutine calc_twostate_model1(moded,phres,addrec,lokph,mc,ceq)
! subroutine calc_twostate_modelny(moded,phres,addrec,lokph,mc,ceq)
! The routine _john works OK but I am testing a modification
! moded is 0, 1 or 2 if no, first or 2nd order derivatives should be calculated
! addrec is addition record
! phres is phase_varres record
! lokph is phase location
! mc is number of constitution fractions
! ceq is current equilibrium record
  implicit none
  integer moded,lokph,mc
  TYPE(GTP_PHASE_ADD), pointer :: addrec
  TYPE(GTP_PHASE_VARRES), pointer :: phres
  TYPE(GTP_EQILIBRIUM_DATA), pointer :: ceq
-----
  subroutine calc_twostate_model_old(moded,phres,addrec,lokph,mc,ceq)
! Failed attempt to decrease the hump when the g2 parameter changes sign
! moded is 0, 1 or 2 if no, first or 2nd order derivatives should be calculated
! addrec is addition record
! phres is phase_varres record
! lokph is phase location
! mc is number of constitution fractions
! ceq is current equilibrium record
  implicit none
  integer moded,lokph,mc
  TYPE(GTP_PHASE_ADD), pointer :: addrec
  TYPE(GTP_PHASE_VARRES), pointer :: phres
  TYPE(GTP_EQILIBRIUM_DATA), pointer :: ceq

```

8.14.11 Debye heat capacity model

The Debye model has been abandoned because the Einstein model is much simpler and both need additional T terms to fit experimental data.

Unfinished.

```
subroutine create_debyecp(addrac)
! enters a record for the debye model
  implicit none
  type(gtp_phase_add), pointer :: addrac
-----
subroutine calc_debyecp(moded,phres,lokadd,lokph,mc,ceq)
! calculates Mauro Debye contribution
! NOTE: values for function not saved, should be done to save calculation time.
! moded: integer, 0=only G, S, Cp; 1=G and dG/dy; 2=Gm dG/dy and d2G/dy2
! phres: pointer, to phase\_varres record
! lokadd: pointer, to addition record
! lokph: integer, phase record
! mc: integer, number of constituents
! ceq: pointer, to gtp\_equilibrium\_data
  implicit none
  integer moded,lokph,mc
  TYPE(gtp\_equilibrium\_data), pointer :: ceq
  TYPE(gtp\_phase\_add), pointer :: lokadd
  TYPE(gtp\_phase\_varres) :: phres
```

8.14.12 Handling diffusion data

This is intended to convert mobility data to diffusion coefficients. using the second derivatives of the Gibbs energy.

```
subroutine create_diffusion(addrac,lokph,text)
  implicit none
  integer lokph
  character text*(*)
  type(gtp_phase_add), pointer :: addrac
-----
subroutine diffusion_onoff(phasetup,bitval)
! switches the bit which calculates diffusion coefficients on/off
! if bitval 0 calculate is turned on, 1 turn off
  implicit none
  integer bitval
  type(gtp_phasetuple) :: phasetup
-----
```

```

subroutine calc_diffusion(moded,phres,lokadd,lokph,mc,ceq)
! calculates diffusion coefficients
! NOTE: values for function not saved, should be done to save calculation time.
! moded: integer, 0=only G, S, Cp; 1=G and dG/dy; 2=Gm dG/dy and d2G/dy2
! phres: pointer, to phase\_varres record
! lokadd: pointer, to addition record
! lokph: integer, phase record
! mc: integer, number of constituents
! ceq: pointer, to gtp\_equilibrium\_data
  implicit none
  integer moded,lokph,mc
  TYPE(gtp\_equilibrium\_data), pointer :: ceq
  TYPE(gtp\_phase\_add), pointer :: lokadd
  TYPE(gtp\_phase\_varres) :: phres
-----
subroutine get_diffusion_matrix(phtup,mdm,dcval,ceq)
! extracts calculated diffusion coefficients for a phase tuple
! phtup phase tuple
! dcval diffusion matrix
! ceq: pointer, to gtp\_equilibrium\_data
  implicit none
  integer mdm
  double precision dcval(mdm,*)
  TYPE(gtp\_phasetuple) :: phtup
  TYPE(gtp\_equilibrium\_data), pointer :: ceq

```

8.14.13 List additions

When listing data for a phase with addition the relevant information must be listed also for additions.

```

subroutine list_addition(unit,CHTD,phname,ftyp,lokadd)
! list description of an addition for a phase on unit
! used when writing databases files and phase data
  implicit none
  integer unit,ftyp
! CHTD is letter for TDB files TYPE\_DEFINITION ... suck
  character CHTD*1,phname*(*)
  TYPE(gtp\_phase\_add), pointer :: lokadd
-----
subroutine list_addition_values(unit,phres)
! lists calculated values for this addition
! Used for the command CALCULATE PHASE to inform about additions
  implicit none
  integer unit

```



```
TYPE(gtp_phase_varres), pointer :: phres
```

8.15 Calculation

There are many subroutines involved in calculating the Gibbs energy for a system and to retrieve values afterwards. Some are explained in connection with what they calculate, for example the magnetic contribution in 8.14.3.

8.15.1 Calculate for one phase

This subroutine calculates the Gibbs energy and all first and second derivatives with respect to T , P and constituents for the specified phase and composition set using the current values of T , P and constitution of the phase (set by `set_constitution`, see 8.5.1). It also calculates all other properties stored in the property records.

It is possible to calculate only G by setting `moded=0`, only G and first derivatives if `moded=1` and also second derivatives with `moded=2`. This routine calls `calcg_internal` to do the calculations after some checks.

```
subroutine calcg(iph,ics,moded,lokres,ceq)
! calculates G for phase iph and composition set ics in equilibrium ceq
! checks first that phase and composition set exists
! Data taken and stored in equilibrium record ceq
! lokres is set to the phase_varres record with all fractions and results
! moded is 0, 1 or 2 depending on calculating no, first or 2nd derivatives
  implicit none
  TYPE(gtp_equilibrium_data), pointer :: ceq
  integer iph,ics,moded,lokres
```

8.15.2 Model independent routine for one phase calculation

This is the central subroutine to calculate G and derivatives for all kinds of phases. At present only the CEF and ionic liquid model is implemented. It calls many other calculating subroutines, some are described in connection with the property they calculate, like magnetic contribution.

```
subroutine calcg_internal(lokph,moded,cps,ceq)
! Central calculating routine calculating G and everything else for a phase
! ceq is the equilibrium record, cps is the phase_varres record for lokph
! moded is type of calculation, 0=only G, 1 G and first derivatives
! 2=G and all second derivatives
! Can also handle the ionic liquid model now ....
  implicit none
  integer lokph,moded
  TYPE(gtp_equilibrium_data), pointer :: ceq
  TYPE(gtp_phase_varres), target :: cps
```

8.15.3 A utility routine

This is used when a phase has permutations, see 5.6.1

```
subroutine setendmemarr(lokph,ceq)
! stores the pointers to all ordered and disordered endmemners in arrays
! intended to allow parallel calculation of parameters
! UNUSED ??
  implicit none
  integer lokph
  TYPE(gtp_equilibrium_data), pointer :: ceq
```

8.15.4 Calculate and list results for one phase

This is mainly a debugging routine that calculates and lists for a specific phase the Gibbs energy and all first and second derivatives by calling calcg using the current values of T , P and constitution. It does not iterate and can thus not calculate an equilibrium.

```
subroutine tabder(ip,ics,times,ceq)
! tabulate derivatives of phase ip with current constitution and T and P
  implicit none
  integer ip,ics,times
  TYPE(gtp_equilibrium_data), pointer :: ceq
```

8.15.5 Calculate an interaction parameter

This is called by calcg_internal to calculate the value of an interaction parameter and its derivatives and add this to all property arrays.

```
subroutine cgint(lokph,lokpty,moded,vals,dvals,d2vals,gz,ceq)
! calculates an excess parameter that can be composition dependent
! gz%yfirem are the site fractions in the end member record
! gz%yfrint are the site fractions in the interaction record(s)
! lokpty is the property index, lokph is the phase record
! moded=0 means only G, =1 G and dG/dy, =2 all
  implicit none
  integer moded,lokph
  TYPE(gtp_property), pointer :: lokpty
  TYPE(gtp_parcalc) :: gz
  double precision vals(6),dvals(3,gz%nofc)
  TYPE(gtp_equilibrium_data), pointer :: ceq
```

8.15.6 Calculate ideal configurational entropy

This calculates the ideal configurational entropy summed over all sublattices.

```
subroutine config_entropy(moded,ns1,nkl,phvar,tval)
! calculates configurational entropy/R for phase lokph
  implicit none
  integer moded,ns1
  integer, dimension(ns1) :: nkl
  TYPE(gtp_phase_varres), pointer :: phvar
```

8.15.7 Calculate ionic liquid configurational entropy

The ionic liquid model assumes ideal mixing on each sublattice but the site ratios are not constant.

```
subroutine config_entropy_i2sl(moded,ns1,nkl,phvar,i2slx,tval)
! calculates configurational entropy/R for ionic liquid model
! Always 2 sublattices, the sites depend on composition
!  $P = \sum_j (-v_j) y_j + Q y_{Va}$ 
!  $Q = \sum_i v_i y_i$ 
! where v is the charge on the ions. P and Q calculated by set_constitution
  implicit none
  integer moded,ns1,i2slx(2)
  integer, dimension(ns1) :: nkl
  TYPE(gtp_phase_varres), pointer :: phvar
```

8.15.8 The configurational entropy for other models

This is for polymers consisting of long chains that can be can have very different molar volumes.

```
subroutine config_entropy_cqc_classicqc(moded,ncon,phvar,phrec,tval)
!
! calculates configurational entropy/R for the classical quasichemical liquid
!
! THIS WORKS OK 2019-01-10: DO NOT CHANGE ANYTHING!! works for qcmodel=1
! routine for qcmodel=2 and 3 laret
!
! only question is the parameter, value of  $G = K \cdot T \cdot R / 2$ ;
!  $K=-10$ ,  $T=600$  means  $G = -10 \cdot 600 \cdot R / 2 = -3000 \cdot R$  gives same curves as in paper.
!
! moded=0 only G, =1 G and dG/dy, =2 G, dG/dy and d2G/dy1/dy2
! ncon is number of constituents
! phvar is pointer to phase_varres record
! phrec is the phase record
```

```

! tval is current value of T
  implicit none
  integer moded,ncon
  TYPE(gtp_phase_varres), pointer :: phvar
  TYPE(gtp_phaserecord) :: phrec
  double precision tval
-----
  subroutine config_entropy_floryhuggins(moded,nofc,phvar,tval,fhv,dfhv,d2fhv)
! calculates configurational entropy/R for a phase with Flory-Huggins model
! moded=0 only G; =1 G and dG/dy; =2; G, dG/dy and d2G/dy2
! nofc number of constituents, phvar phase_varres record
  implicit none
  integer moded,nofc
  TYPE(gtp_phase_varres), pointer :: phvar
! fvh(1,1) is FH volume for constituent 1 etc.
  double precision tval,fhv(nofc,*),dfhv(nofc,3,*),d2fhv(nofc,*)
-----
  subroutine config_entropy_cqc6(moded,ncon,phvar,phrec,tval)
!
! calculates configurational entropy/R for the corrected quasichemical liquid
! Rewritten 2019-01-12 based on cqc-classicqc which seems correct
! test1: qcmodel=1: OK for zhalf=1 and 3
! test2: qcmodel=2: OK!!
! test3: qcmodel=3: OK for SR0, problems for miscibility gap
!
! only question is the parameter, value of  $G = K \cdot T \cdot R / 2$ ;
!  $K=-10$ ,  $T=600$  means  $G = -10 \cdot 600 \cdot R / 2 = -3000 \cdot R$  gives same curves as in paper.
!
! moded=0 only G, =1 G and dG/dy, =2 G, dG/dy and d2G/dy1/dy2
! ncon is number of constituents
! phvar is pointer to phase_varres record
! phrec is the phase record
! tval is current value of T
  implicit none
  integer moded,ncon
  TYPE(gtp_phase_varres), pointer :: phvar
  TYPE(gtp_phaserecord) :: phrec
  double precision tval

```

8.15.9 Push/pop constituent fraction product on stack

These subroutines are used to push/pop current values of the product of constituent fractions and its derivatives before calculating an interaction parameter.

```

subroutine push_pyval(pystack,intrec,pmq,pyq,dpyq,d2pyq,moded,iz)

```

```

! push data when entering an interaction record
  implicit none
  integer pmq,moded,iz
  double precision pyq,dpyq(iz),d2pyq(iz*(iz+1)/2)
  type(gtp_pystack), pointer :: pystack
  type(gtp_interaction), pointer :: intrec
-----
  subroutine pop_pyval(pystack,intrec,pmq,pyq,dpyq,d2pyq,moded,iz)
! pop data when entering an interaction record
  implicit none
  integer iz,pmq,moded
  double precision pyq,dpyq(iz),d2pyq(iz*(iz+1)/2)
  type(gtp_pystack), pointer :: pystack
  type(gtp_interaction), pointer :: intrec

```

8.15.10 Calculate disordered fractions from constituent fractions

This is used when there are several fraction sets of a phase. The values of the second fraction set (also called the disordered fraction set) is calculated by this subroutine. These disordered fractions can be used to calculate a “disordered” part of the Gibbs energy with its own set of parameters.

```

  subroutine calc_disfrac(lokph,lokcs,ceq)
! calculate and set disordered set of fractions from sitefractions
! The first derivatives are dxidyj. There are no second derivatives
! TYPE(gtp_fraction_set), pointer :: disrec
  implicit none
  integer lokph,lokcs
  TYPE(gtp_equilibrium_data), pointer :: ceq
-----
  subroutine calc_disfrac2(phord,phdis,ceq)
! calculate and set disordered set of fractions from sitefractions
! The first derivatives are dxidyj. There are no second derivatives
! TYPE(gtp_fraction_set), pointer :: disrec
  implicit none
  TYPE(gtp_phase_varres), target :: phord
  TYPE(gtp_phase_varres), pointer :: phdis
  TYPE(gtp_equilibrium_data), pointer :: ceq

```

8.15.11 Disorder constituent fractions

The Gibbs energy for the “partitioned” phases, like FCC and BCC which can have order/disorder transformations, see 2.5, the “ordered part” is calculated twice, once with the original constituent fractions and once with these set equal to their disordered value. The reason for this is that the “disordered part” should be complete i.e. include also the disordered part of the “ordered part” as the disordered partitions.

This is now an optional way to handle partitioning, the recommended way is to simply add the Gibbs energy is simply added from the two fraction sets. This subroutine sets the fractions to their disordered values.

```

subroutine disordery(phvar,ceq)
! sets the ordered site fractions in FCC and other order/disordered phases
! equal to their disordered value in order to calculate and subtract this part
! phvar is pointer to phase_varres for ordered fractions
  implicit none
  TYPE(gtp_phase_varres), pointer :: phvar
  TYPE(gtp_equilibrium_data), pointer :: ceq
-----
subroutine disordery2(lokdc,phvar,disrec,ceq)
! subroutine disordery2(phdis,phvar,disrec,ceq)
! sets the ordered site fractions in FCC and other order/disordered phases
! equal to their disordered value in order to calculate and subtract this part
! phvar is pointer to phase_varres for ordered fractions
! phdis is pointer to phase_varres for disordered fractions
  implicit none
  TYPE(gtp_phase_varres), pointer :: phvar
  TYPE(gtp_fraction_set), pointer :: disrec
  TYPE(gtp_equilibrium_data), pointer :: ceq

```

8.15.12 Subroutine to calculate the UNIQUAC model

A routine to calculate configurational entropy and residual energy for the UNIQUAC model. It seems to work! Documentation of the model is missing (except comments inside this subroutine) but there is a macro file, “uniquac.OCM”, that explains how to enter species and calculate binary and ternary diagrams. There are several parts of this documentation describing how bits, species etc. has been been changed to accomodate this model.

```

subroutine uniquac_model(moded,ncon,phres,ceq)
! Calculate the Gibbs energy of the UNIQUAC model (Abrams et al 1975)
! Modified 2018/Oct, Nov, Dec
! It returns UNIQUAC G and first and second derivatives of G in phres%gval etc.
! The values of q_i and r_i are be stored in species record, not identifiers
! The residual term should be stored as a UQTAU identifier
  implicit none
  integer moded,ncon
  TYPE(gtp_equilibrium_data), pointer :: ceq
  TYPE(gtp_phase_varres), pointer :: phres

```

8.15.13 Set driving force for a phase explicitly

A failed attempt to handle convergence problems.

```

subroutine set_driving_force(iph,ics,dgm,ceq)
! set the driving force of a phase explicitly
  implicit none
  type(gtp_equilibrium_data), pointer :: ceq
  integer iph,ics
  double precision dgm

```

8.15.14 Extract mass balance conditions

This is used in global grid minimization to extract the set of mass balance conditions. If the current conditions are not all mass balance there is an error return, otherwise the conditions of T and P, the total number of moles and the mole fractions of all components are returned.

```

subroutine extract_massbalcond(tpval,xknown,antot,ceq)
! extract T, P, mol fractions of all components and total number of moles
! for use when minimizing G for a closed system. Probably redundant
  implicit none
  double precision, dimension(*) :: tpval,xknown
  double precision antot
  TYPE(gtp_equilibrium_data), pointer :: ceq

```

8.15.15 Saving and restoring a phase constitution

These are used during step and map to help handling convergence problems

```

subroutine save_constitutions(ceq,copyofconst)
! copy the current phase amounts and constitutions to be restored
! if calculations fails during step/map
! DANGEROUS IF NEW COMPOSITION SETS CREATED
  implicit none
  TYPE(gtp_equilibrium_data), pointer :: ceq
  double precision, allocatable, dimension(:) :: copyofconst
-----
subroutine restore_constitutions(ceq,copyofconst)
! restore the phase amounts and constitutions from copyofconst
! if calculations fails during step/map
! DANGEROUS IF NEW COMPOSITION SETS CREATED
  implicit none
  TYPE(gtp_equilibrium_data), pointer :: ceq
  double precision copyofconst(*)

```

8.16 Grid minimizer

When there is only mass balance conditions it is possible to use a global minimization technique to find start points for a more standard Newton-Raphson technique. The latter has the disadvantage that it can only find local minimas.

In the global minimization all solution phases are divided into a grid with fixed compositions and the Gibbs energy for each gridpoint is calculated. These gridpoints are then searched until one finds a set that encloses the given overall composition which represent the minimal Gibbs energy. It is thus necessary to know the overall composition.

The phases the gridpoints in the solution are then identified and possibly one can merge gridpoints in the same phase unless they are separated by a miscibility gap.

Some care should be taken with ordered phases as they have several identical sublattices and one thus can reduce significantly the number of gridpoints generated if this is taken into account.

8.16.1 Global Gridminimizer

This is the main grid minimizing subroutine. It can be called also after an equilibrium calculation, see section 8.16.8, to check if there are any phases with a gridpoint below the calculated G hyper-surface. That is useful if the conditions does not allow, for example if T is not known, that the gridminimizer is called before the equilibrium calculation.

```
subroutine global_gridmin(what,tp,xknown,nvsph,iph1,icsl,aph1,&
    nyph1,cmu,ceq)
!
! Starting rewriting 2017-02-01
!
! finds a set of phases that is a global start point for an equilibrium
! calculation at T and P values in tp and known mole fraction in xknown
! It is intentional that this routine is independent of current conditions
! It returns: nvsph stable phases, list of phases in iph1, amounts in aph1,
! nyph1(i) is redundant, cmu are element chemical potentials of solution
! WHAT determine what to do with the results, 0=just return solution,
! 1=enter stable set and constitution of all phases in gtp datastructure
! and create composition sets if necessary (and allowed)
! what=-1 will check if any gridpoint is below current calculated equilibrium
! ?? removed what -1 170428/BoS
    implicit none
! nyph1(j) is the start position of the constituent fractions of phase j in
    integer, dimension(*) :: iph1,nyph1,icsl
    integer what,nvsph

    TYPE(gtp_equilibrium_data), pointer :: ceq
    double precision, dimension(2) :: tp
! cmu(1..nrel) is the chemical potentials of the solution
```



```
double precision, dimension(*) :: xknown,aphl,cmu
```

8.16.2 Generate grid

There are several different grid generators depending on the type of phase for example ionic or with order/disorder transformation. It is also a possibility to generate a denser grid.

The ordered fcc, bcc and hcp with 4 sublattice model can have significantly reduced number of gridpoints. This is not yet implemented. For the ordered 4 sublattice model we can exclude all permutations of the endmembers, for example only include A:A:A:B but not A:A:B:A, A:B:A:A and B:A:A:A. This reduces the number of gridpoints that must be calculated.

The same when a phase have charged constituents with many anemembers that have net charge. This is implemented but not optimized.

```
subroutine generate_grid(mode,iph,ngg,nrel,xarr,garr,ny,yarr,gmax,ceq)
! Different action depending of the value of mode,
! for mode<0: (will no longer be used ... )
!   return the number of gridpoints that will be generated for phase iph in ngg
! for mode=0:
!   on entry ngg is dimension of garr
!   on exit ngg is number of generated gridpoints ...
!   return garr(i) gibbs energy and xarr(1,i) the compositions of gridpoint i
! for mode>0:
!   return site fractions of gridpoint mode in yarr, number of fractions in ny
!   iph is phase number, ngg is number of gridpoints, nrel number of elements,
! if mode=0:
!   return xarr mole fractions of gridpoints, garr Gibbs energy of gridpoints,
!   ngg is dimension of garr, gmax maximum G (start value for chem.pot)
! if mode>0:
!   "mode" is a gridpoint of this phase in solution, return number of
!   constituent fractions in ny and fractions in yarr for this gridpoint
! The current constitution is restored at the end of the subroutine
  implicit none
  TYPE(gtp_equilibrium_data), pointer :: ceq
  integer mode,iph,ngg,nrel,ny
  real xarr(nrel,*),garr(*)
  double precision yarr(*),gmax
-----
subroutine generic_grid_generator(mode,iph,ngg,nrel,xarr,garr,ny,yarr,gmax,ceq)
! This generates grid for any phase
! mode=0 generate grid for phase iph and mole fraction and G for all points
!   ngg on entry is max number of gridpoints, on exit number of gridpoints
!   nrel is number of elements
!   xarr(1..nrel,gp) is composition of gripoint gp, garr(iv) its G
!   ny,yarr,gmax not used
```

```

! mode>0 return constitution for gridpoint number mode in yarr
!   iph is returned as phase index for gripoint mode
!   xarr(1..nrel) the composition at the gripoint, garr not nused
!   ny is number of constituent fractions
!   yarr are the constituent fractions
!   gmax not used ??
!
  implicit none
  integer mode,iph,ngg,nrel,ny
  real xarr(nrel,*),garr(*)
  double precision yarr(*),gmax
  type(gtp_equilibrium_data), pointer :: ceq
-----
  subroutine generate_dense_grid(mode,iph,ngg,nrel,xarr,garr,ny,yarr,gmax,ceq)
! generates more gridpoints than default generate_grid
! Different action depending of the value of mode,
! for mode<0:
!   return the number of gridpoints that will be generated for phase iph in ngg
! for mode=0:
!   return garr(i) gibbs energy and xarr(1,i) the compositions of gridpoint i
! for mode>0:
!   return site fractions of gridpoint mode in yarr, number of fractions in ny
!   iph is phase number, ngg is number of gridpoints, nrel number of elements,
! if mode=0:
!   return xarr mole fractions of gridpoints, garr Gibbs energy of gridpoints,
!   ngg is dimension of garr
! if mode>0:
!   "mode" is a gridpoint of this phase in solution, return number of
!   constituent fractions in ny and fractions in yarr for this gridpoint
! The current constitution is restored at the end of the subroutine
  implicit none
  TYPE(gtp_equilibrium_data), pointer :: ceq
  integer mode,iph,ngg,nrel,ny
  real xarr(nrel,*),garr(*)
  double precision yarr(*),gmax
-----
  subroutine generate_ionliq_grid(mode,iph,ngg,nrel,xarr,garr,ny,yarr,gmax,ceq)
! generates gridpoints for ionic liquid (also dense)
! Different action depending of the value of mode,
! for mode<0:
!   return the number of gridpoints that will be generated for phase iph in ngg
! for mode=0:
!   return garr(i) gibbs energy and xarr(1,i) the compositions of gridpoint i
! for mode>0:
!   return site fractions of gridpoint mode in yarr, number of fractions in ny
!   iph is phase number, ngg is number of gridpoints, nrel number of elements,

```

```

! if mode=0:
!   return xarr mole fractions of gridpoints, garr Gibbs energy of gridpoints,
!   ngg is dimension of garr
! if mode>0:
!   "mode" is a gridpoint of this phase in solution, return number of
!   constituent fractions in ny and fractions in yarr for this gridpoint
! The current constitution is restored at the end of the subroutine
  implicit none
  TYPE(gtp_equilibrium_data), pointer :: ceq
  integer mode,iph,ngg,nrel,ny
  real xarr(nrel,*),garr(*)
  double precision yarr(*),gmax
-----
  subroutine generate_fccord_grid(mode,iph,ngg,nrel,xarr,garr,ny,yarr,gmax,ceq)
! This generates grid for a phase with 4 sublattice fcc/bcc/hcp ordering
! NO LONGER USED: mode<0 just number of gridpoints in ngg, for allocations
! mode=0 calculate mole fraction and G for all gridpoints
! mode>0 return constitution for gridpoint mode in yarr
  implicit none
  integer mode,iph,ngg,nrel,ny
  real xarr(nrel,*),garr(*)
  double precision yarr(*),gmax
  type(gtp_equilibrium_data), pointer :: ceq
-----
  subroutine generate_charged_grid(mode,iph,ngg,nrel,xarr,garr,ny,yarr,gmax,ceq)
! This generates grid for a phase with charged constituents
! mode<0 just number of gridpoints in ngg, needed for allocations
! mode=0 calculate mole fraction and G for all gridpoints
! mode>0 return constitution for gridpoint mode in yarr
  implicit none
  integer mode,iph,ngg,nrel,ny
  real xarr(nrel,*),garr(*)
  double precision yarr(*),gmax
  type(gtp_equilibrium_data), pointer :: ceq

```

8.16.3 Calculate gridpoint

The Gibbs energy for each gridpoint is calculated as well as its composition.

```

subroutine calc_gridpoint(iph,yfra,nrel,xarr,gval,ceq)
! called by global minimization routine
! Not adopted to charged crystalline phases as gridpoints have net charge
! but charged gripoints have high energy, better to look for neutral ones ...
  implicit none
  real xarr(*),gval

```

```

integer iph,nrel
double precision yfra(*)
TYPE(gtp_equilibrium_data), pointer :: ceq

```

8.16.4 Calculate endmember

These subroutines are used by the calc_gridpoint routine and also to calculate a user defined reference state..

```

subroutine calcg_endmember(iphx,endmember,gval,ceq)
! calculates G for one mole of real atoms for a single end member
! used for reference states. Restores current composition (but not G or deriv)
! endmember contains indices in the constituent array, not species index
! one for each sublattice
  implicit none
  integer iphx
  double precision gval
  integer endmember(maxsubl)
  TYPE(gtp_equilibrium_data), pointer :: ceq
-----
subroutine calcg_endmemberx(iphx,endmember,gval,ceq)
! calculates G for single end member with current number of atoms
! used for reference states. Restores current composition (but not G or deriv)
! endmember contains indices in the constituent array, not species index
! one for each sublattice
  implicit none
  integer iphx
  double precision gval
  integer endmember(maxsubl)
  TYPE(gtp_equilibrium_data), pointer :: ceq
-----
subroutine calcg_endmember6(iph,endmember,gval,ceq)
! calculates G AND ALL DERIVATEVS wrt T and P for one mole of real atoms
! for a single end member, used for reference states.
! Restores current composition and G (but not deriv)
! endmember contains indices in the constituent array, not species index
! one for each sublattice
  implicit none
  integer iph
  double precision gval(6)
  integer endmember(maxsubl)
  TYPE(gtp_equilibrium_data), pointer :: ceq

```

8.16.5 Calculate minimum of grid

The search starts from the lowest G value for the pure elements. (This seems obvious but makes it impossible to have a grid for only fcc with carbon dissolved as fcc does not exist for pure carbon). The selected gridpoints always represent a hyperplane (number of dimensions equal to that of the number of components) of Gibbs energy over the whole composition range. Then the whole grid is searched for the gridpoint that has the most negative deviation from this hyperplane. Then one gridpoint in the existing hyperplane is replaced by this in such a way that the overall mass balance is inside the points forming the plane. Then a new search is made and a point is replaced until there are no points below the hyperplane. The points forming this plane represent the solution. There will always be as many points as components.

```
subroutine find_gridmin(kp,nrel,xarr,garr,xknown,jgrid,phfrac,cmu,trace)
! there are kp gridpoints, nrel is number of components
! composition of each gridpoint in xarr, G in garr
! xknown is the known overall composition
! return the gridpoints of the solution in jgrid, the phase fraction in phfrac
! cmu are the final chemical potentials
  implicit none
  integer, parameter :: jerr=50
  integer kp,nrel
  integer, dimension(*) :: jgrid
  real xarr(nrel,*),garr(*)
  double precision xknown(*),phfrac(*),cmu(nrel)
  logical trace
```

8.16.6 Merge gridpoints in same phase

This subroutine tries to check if the number of gridpoints can be reduced by merging gridpoints in the same phase. Care must be taken that there can be miscibility gaps between points.

This subroutine may automatically create new composition sets if necessary (unless the user has set the appropriate bit to prevent that).

```
subroutine merge_gridpoints(nv,iphl,aphl,nyphl,yphl,trace,nrel,xsol,cmu,ceq)
!
! BEWARE not adopted for parallel processing
!
! if the same phase has several gridpoints check if they are really separate
! (miscibility gaps) or if they can be merged. Compare them two by two
! nv is the number of phases, iphl(i) is the index of phase i, apha(i) is the
! amount of phase i, nyphl is the number of site fractions for phase i,
! and yphl is the site fractions packed together
!
  implicit none
```

```

TYPE(gtp_equilibrium_data), pointer :: ceq
integer nv,nrel
integer, dimension(*) :: iphl,nyphl
double precision, dimension(*) :: aphl,yphl,cmu
logical trace
real xsol(maxel,*)

```

8.16.7 Set constitution of metastable phases

Phases not part of the final solution will have their constitution set to a gridpoint that is closest to the final hyperplane.

```

subroutine set_metastable_constitutions2(pph,nrel,nphl,iphx,xarr,garr,&
    nr,iphl,cmu,ceq)
! this subroutine goes through all the metastable phases
! after a global minimization and sets the constitution to the most
! favourable one. Later care should be taken that exiting higher composition
! sets are not set equal to the stable
! pph  number of phases for which a grid has been calculated
! nrel number of components
! nphl(p) is last gridpoint for phase(p), nphl(0)=0, p=1,pph
! iphx(p) phase number of phase(p) (skipping dormant and suspended phases)
! xarr(1..nrel,i) composition of gridpoint i
! garr(i) Gibbs energy/RT for gridpoint i
! nr    is the number of stable phases in the solution
! iphl(s) the phase number of the stable phases s (not ordered)
! cmu   are the chemical potentials/RT of the solution
! ceq   equilibrium record
! called by global_gridmin
implicit none
integer pph,nrel,nr
integer, dimension(0:*) :: nphl
integer, dimension(*) :: iphl,iphx
double precision, dimension(*) :: cmu
real garr(*),xarr(nrel,*)
type(gtp_equilibrium_data), pointer :: ceq

```

8.16.8 Check of equilibrium using grid minimizer

This subroutine can be used after an equilibrium calculation when the conditions does not allow the grid minimizer to be used (if T, P are not known or all conditions are not mass balance). It calculates the same grid as the global gridmin and then checks if any gripoint is below the current hyperplane definde by the calculated chemical potentials. If so it reports the phase and composition. It does not correct the equilibrium.

```

logical function global_equil_check1(mode,addtuple,yfr,ceq)
! subroutine global_equil_check(ceq,newceq)
!
! THIS MUST BE REWRITTEN
!
! This subroutine checks there are any gridpoints below the calculated solution
! if not it is taken as a correct global equilibrium
! This avoids creating any new composition sets but may fail in some cases
! to detect that the equilibrium is not global.
! mode=1 means try to recalculate equilibrium if not global (not implemented)
! if a gridpoint below is found addtuple and yfr returned with this
  implicit none
  integer mode,addtuple
  double precision, allocatable, dimension(:) :: yfr
  TYPE(gtp_equilibrium_data), pointer :: ceq

```

8.16.9 Separate composition set constitutions that has merged

During step and map the constitution of two composition sets may merge above the top of the miscibility gap. This routine will check for that and separate them if that has happend.

```

subroutine separate_constitutions(ceq)
! Go through all entered phases and if there are two composition sets
! that have similar constitutions then separate them
! Used during mapping of for example Fe-Cr to detect the miscibility gap
  implicit none
  TYPE(gtp_equilibrium_data), pointer :: ceq

```

8.16.10 Check if two phases are allotropes

Check if two phases are allotropes, i.e. have identical models and composition.

```

logical function allotropes(irem,iadd,iter,ceq)
! This function return TRUE if the phases indicated by IREM and IADD both have
! fixed and identical composition, i.e. they are compounds and allotropes
! Such a transition can cause problems during a STEP command.
  implicit none
  TYPE(gtp_equilibrium_data), pointer :: ceq
  integer iadd,irem,iter
-----
logical function same_stoik(iph,jph)
! MAYBE IDENTICAL TO ALLOTROPEs ?
! return TRUE if phase iph and jph are both stoichiometric and have the
! same composition Used to check when adding a phase during equilibrium

```

```

! calculation as it normally fails to have two such phases stable
  implicit none
  integer iph,jph
-----
  logical function fixedcomposition(iph)
! returns TRUE if phase cannot vary its composition
  integer iph

```

8.17 Miscellaneous things

Things that does not fit anywhere else.

8.17.1 Phase record location

```

integer function phvarlok(lokph)
! return index of the first phase_varres record for phase with location lokph
! needed for external routines as phlista is private
  implicit none
  integer lokph

```

8.17.2 Numbers an interaction tree for permutations

For permutations it turned out to be necessary to keep track of the individual interaction records to know the permutation. This subroutine indexes the interaction records for each endmember of a phase.

```

subroutine palmtree(lokph)
! Initiates a numbering of all interaction trees of an endmember of a phase
  implicit none
  integer lokph

```

8.17.3 Sorting constituent fractions

```

subroutine sortinphtup(n,m,xx)
! subroutine to sort the values in xx which are in phase and compset order
! in phase tuple order. This is needed by the TQ interface
! The number of values belonging to the phase is m (for example composition)
! argument ceq added as new composition sets can be created ...
  integer n,m
!   double precision xx(n*m)
  double precision xx(*)
!   type(gtp_equilibrium_data), pointer :: ceq

```


8.17.4 Get model parameter identifier index

Extract the model parameter index.

```
integer function get_mpi_index(mpi)
! Return the index of a model parameter identifier
character mpi*(*)
-----
integer function getmqindex()
! This is necessary because mqindex is private, replaced by getmpiindex ...
```

8.17.5 Check that certain things are allowed

To have a uniform check if something is allowed. One must not enter a phase before there are any elements for example.

```
logical function allowenter(mode)
! Check if certain commands are allowed
! mode=1 means entering an element or species
! mode=2 means entering a phase
! mode=3 means entering an equilibrium
! returns TRUE if command can be executed
implicit none
integer mode
```

8.17.6 Check proper symbol

There are many symbols and names in the GTP package. In general a symbol must start with a letter A-Z. All symbols are also case insensitive, i.e. upper and lower case are treated as the same. Most symbols may contain digits as second and later character. Some symbols and names may contain special characters like “_” and others.

This subroutine checks this.

```
logical function proper_symbol_name(name,typ)
! checks that name is a proper name for a symbol
! A proper name must start with a letter A-Z
! for typ=0 it must contain only letters, digits and underscore
! for typ=1 it may contain also +, - maybe ?
! It must not be equal to a state variable
implicit none
integer typ
character name*(*)
```

8.17.7 List things for debugging

List constitutions in some order or other.

```
subroutine compmassbug(ceq)
! debug subroutine
  type(gtp_equilibrium_data), pointer :: ceq
-----
subroutine list_free_lists(lut)
! for debugging the free lists and routines using them
  implicit none
  integer lut
```

8.17.8 The amount of a phase is set to a value

I am not sure when this is used or needed.

```
subroutine set_phase_amounts(jph,ics,val,ceq)
! set the amount formula units of a phase. Called from user i/f
! iph can be -1 meaning all phases, all composition sets
  implicit none
  integer jph,ics
  double precision val
  TYPE(gtp_equilibrium_data), pointer :: ceq
```

8.17.9 Set the default constitution of a phase

I am not sure when this is used and if it is needed.

```
subroutine set_default_constitution(iph,ics,ceq)
! the current constitution of (iph#ics) is set to its default constitution
! (if any), otherwise a random value. The amount of the phase not changed
  implicit none
  integer iph,ics
  TYPE(gtp_equilibrium_data), pointer :: ceq
```

8.17.10 Subroutine to prepare for an equilibrium calculation

Unfinished

```
subroutine todo_before(mode,ceq)
! this could be called before an equilibrium calculation
! It should remove any phase amounts and clears CSSTABLE
```

```

! DUMMY
!
  implicit none
  TYPE(gtp_equilibrium_data), pointer :: ceq
  integer mode

```

8.17.11 Subroutine to clean up after an equilibrium calculation

This now checks if there are any composition sets with the AUTO bit set meaning that they have been created by the grid minimizer in this equilibrium calculation. If so it tries to move the stable phases to the lowest possible composition set, taking care that user defined composition sets with a given default constitution is honored, i.e a fcc carbide is not set as composition set 1 if the user has defined a metallic fcc phase with low carbon as the first.

It then removes unstable composition sets with the AUTO bit set and any stable composition sets with the AUTO bit set have this bit cleared.

```

subroutine todo_after_found_equilibrium(mode,addtuple,ceq)
! this is called after an equilibrium calculation by calceq2 and calceq3
! It marks stable phase (set CSSTABLE and remove any CSAUTO)
! It removes redundant unstable composition sets created automatically
! (CSAUTO set). It will also shift stable composition sets to lowest
! possible (it will take into account if there are default constituent
! fractions, CSDEFCON set).
! mode determine some of the actions, at present only >0 or <0 matters
!
! >>>>>>>>> THIS IS DANGEROUS IN PARALLEL PROCESSING
! It should work in step and map as a composition set that once been stable
! will never be removed except if one does global minimization during the
! step and map. The function global_equil_check works on a copy of the
! ceq record and creates only a grid, it does not create any composition sets.
! NOTE that automatically entered metallic-FCC and MC-carbides may shift
! composition sets. Such shifts can be avoided by manual entering composition
! sets with default constitutions, but that does not always work as comparing
! a stable constitution with several defaults is not trivial ...
!
  implicit none
  TYPE(gtp_equilibrium_data), pointer :: ceq
  integer mode,addtuple

```

8.17.12 Select composition set for stable phase

After an equilibrium calculation there may be automatically created composition sets by the grid-minimizer that are not needed. These routines tries to remove unneeded sets and also shift the used ones to the lowest composition set. It tries to take into account the default constitutions for

user defined composition sets for example an fcc#1 with a small amount of C and fcc#2 with a large amount of C. Then a carbide rich fcc phase should be fcc#2 and not fcc#1 even if fcc#1 is not stable.

```

subroutine checkdefcon(lokics,lokjcs,fit,ceq)
! check if composition of lokics fits default constitution in lokjcs
! return TRUE if lokics fits default in lokjcs
! NOTE lokics and lokjcs can be the same!!
  integer lokics,lokjcs,fit
  type(gtp_equilibrium_data), pointer :: ceq
-----
subroutine shiftcompsets(ceq)
! check phase with several composition sets if they should be shifted
! to fit the default constitution better
! IGNORE UNSTABLE COMP.SETS
  type(gtp_equilibrium_data), pointer :: ceq
-----
subroutine switch_compsets2(lokph,ics1,ics2,ceq)
! copy constitution and results from ic2 to ic1 and vice versa
  integer lokph,ics1,ics2
  type(gtp_equilibrium_data), pointer :: ceq

```

8.17.13 Select composition set for stable phase, maybe not used

```

subroutine select_composition_set(iph,ics,yarr,ceq)
! PROBABLY NOT USED but should be implemented
! if phase iph wants to become stable and there are several user defined
! composition sets with default composition limits this subroutine tries to
! select the one that fits these limits best.
! For example if an FCC phase that could be an austenite (low carbon content)
! or a cubic carbo-nitride (high carbon or nitrogen content, low vacancy)
! Less easy to handle ordered phases like B2 or L1_2 as ordering can be
! in any sublatittice ... but with option B and F that is possible
  implicit none
  TYPE(gtp_equilibrium_data), pointer :: ceq
  double precision, dimension(*) :: yarr
  integer iph,ics

```

8.17.14 Check consitency of phase varres

```

subroutine verify_phase_varres_array(ieq,verbose)
! This subroutine checks that the phase varres array is consistent
! in equilibrium ieq. For ieq=1 it also checks the free list
! UNFINISHED and not yet used BUT IMPORTANT
  implicit none

```

```
integer ieq,verbose
```

8.17.15 Not documented

```
subroutine set_emergency_startpoint(mode,phl,amfu,ceq)
! this is called if no previous equilibrium and if grid minimizer
! cannot be used. Select for each element a phase with as much of that
! element as possible to set as stable. Set the remaining phases to a default
! composition. It will never create any composition sets
!
  implicit none
  integer mode,phl(*)
  double precision amfu(*)
  TYPE(gtp_equilibrium_data), pointer :: ceq
-----
  logical function ocv()
! returns TRUE if GSVERBOSE bit is set
-----
  integer function ceqsize(ceq)
! calculates the size in words (4 bytes) of an equilibrium record
  implicit none
  type(gtp_equilibrium_data), pointer :: ceq
-----
  integer function vssize(varres)
! calculates the size in words (4 bytes) of a phase_varres record
  implicit none
  type(gtp_phase_varres) :: varres
```

8.18 Unfinished things

There are many unfinished parts among the other sections, in particular the save/read section.

8.19 TP functions

TP functions are a sub package inside GTP. These can store simple expressions depending on T and P including unary functions LN and EXP (and maybe more). They can return a function value and the first and second derivative with respect to T and P. There are three datatypes associated with these functions described in section 4.

8.19.1 Initiate the TP fun package

```
subroutine tpfun_init(nf,tpres)
! allocate tpfuns and create a free list inside the tpfuns
```

```

    implicit none
    integer nf
! use tpres declared externally for parallel processing
    TYPE(tpfun_parres), dimension(:), allocatable :: tpres

```

8.19.2 Utilities

Because freetpfun is private.

```

    integer function notpf()
! number of tpf functions because freetpfun is private
    implicit none

```

8.19.3 Find a TP function

```

    subroutine find_tpfun_by_name(name,lrot)
! returns the location of a TP function
! if lrot>0 then start after lrot, this is to allow finding with wildcard *
    implicit none
    integer lrot
    character name*(*)
-----
    subroutine find_tpfun_by_name_exact(name,lrot,notent)
! returns the location of a TP function, notent TRUE if not entered
    implicit none
    integer lrot
    logical notent
    character name*(*)

```

8.19.4 Evaluate a TP function

Calculate the value and first and second derivatives.

```

    subroutine eval_tpfun(lrot,tpval,result,tpres)
!     subroutine eval_tpfun(lrot,tpval,symval,result)
! evaluate a TP function with several T ranges
    implicit none
    integer lrot
    double precision tpval(2),result(6),xxx
! changes to avoid memory leak in valgrind
    TYPE(tpfun_parres), dimension(*) :: tpres

```

8.19.5 List the expression of a TP function

```
subroutine list_tpfun(lrot,nosym,str)
! lists a TP symbols with several ranges into string str
! lrot is index of function, if nosym=0 the function name is copied to str
  implicit none
  character str*(*)
  integer nosym,lrot
-----
subroutine list_all_funs(lut)
! list all functions except those starting with _ (parameters)
  implicit none
  integer lut
```

8.19.6 List unentered TP functions

For use when reading from a TDB file as functions may call each other.

```
subroutine list_unentered_funs(lut,nr)
! counts and list functions with TPNOTENT bit set if lut>0
  implicit none
  integer lut,nr
```

8.19.7 Compiles a text string as a TP function

Very messy, requires total rewriting as the whole TP package.

```
subroutine ct1xfn(string,ip,nc,coeff,koder,fromtdb)
!...compiles an expression in string from position ip
!   it can refer to T and P or symbols in fnsym
!   compiled expression returned in coeff and koder
!
! >>> this is very messy
!
!...algorithm for function extraction
! 10*T**2 -5*T*LOG(T) +4*EXP(-5*T**(-1))
!
! AT LABEL 100 start of expression or after (
! sign=1
! -, sign=-1                                goto 200
! +, skip
!
! AT LABEL 200 after sign
! if A-Z                                goto 300
```

```

! if 0-9, extract number          goto 400
! (                               goto 100
! ;                               END or ERROR
! empty                           END or ERROR
! anything else                   ERROR
!
! AT LABEL 300 symbol
! if T or P, extract power if any incl ( ) goto 400
! unary fkn? extract (           goto 100
! symbol                         goto 400
!
! AT LABEL 400 after factor
! -, sign=-1                     goto 200
! sign=1
! +, skip                        goto 200
! )                              goto 400
! ** or ^ extract and store power incl ( ) goto 400
! *                              goto 200
! empty                          goto 900
!
! for TDB compatibility skip #
!
! DO NOT allow unary functions ABOVE(TB) and BELOW(TB)
! check consistency
!   implicit none
!   integer ip,nc,koder(5,*)
!   character string*(*)
!   double precision coeff(*)
!   logical fromtdb
!-----
! subroutine ct1getsym(string,ip,symbol)
! ...extracts an symbol
!   implicit double precision (a-h,o-z)
!   implicit none
!   integer ip
!   character string*(*),symbol*(*)
!-----
! subroutine ct1power(string,ip,ipower)
! ...extracts an integer power possibly surrounded by ( )
!   implicit none
!   integer ip,ipower
!   character string*(*)
!-----
! subroutine ct1mfn(symbol,nranges,tlimits,lokexpr,lrot)
! ...creates a root record with name symbol and temperature ranges
! highest T limit is in tlimits(nranges+1)

```



```

!   implicit double precision (a-h,o-z)
      implicit none
      integer nranges,lrot
      character*(*) symbol
      TYPE(tpfun_expression), dimension(*) :: lokexpr
      real tlimits(*)
-----
      subroutine ct2mfn(symbol,nranges,tlimits,lokexpr,lrot)
!...stores a TPfun in an existing lrot record with name symbol
! and temperature ranges, highest T limit is in tlimits(nranges+1)
      implicit none
      integer nranges,lrot
      character*(*) symbol
      TYPE(tpfun_expression), dimension(*) :: lokexpr
      real tlimits(*)
-----
      subroutine ct1mexpr(nc,coeff,koder,lrot)
!...makes a datastructure of an expression. root is returned in lrot
!   implicit double precision (a-h,o-z)
      implicit none
      integer nc,koder(5,*)
!   TYPE(tpfun_expression), pointer :: lrot
      TYPE(tpfun_expression) :: lrot
!   TYPE(tpfun_expression), pointer :: noexpr
      double precision coeff(*)
-----
      subroutine ct1efn(inrot,tpval,val,tpres)
!...evaluates a datastructure of an expression. Value returned in val
!   inrot is root expression tpfunction record
!   tpval is valuse of T and P, symval is values of symbols
! first and second derivatives of T and P also calculated and returned
! in order F, F.T, F.P, F.T.T, F.T.P, F.P.P
!
! if function already calculated one should never enter this subroutine
!
! It can call "itself" by reference to another TP function and for
! that case one must store results in levels.
      implicit none
      double precision val(6),tpval(*)
      TYPE(tpfun_expression), pointer :: inrot
      TYPE(tpfun_parres), dimension(*) :: tpres

```

8.19.8 Utility to list a TP function

```

subroutine ct1wfn(exprot,tps,string,ip)

```

```

!...writes an expression into string starting at ip
!      lrot is an index to an tpexpr record
!      implicit double precision (a-h,o-z)
!      implicit none
!      character tps(2)*(*)
!      character string*(*)
-----
subroutine ct1wpow(string,ip,tps,mult,npow)
!...writes "ips" with a power if needed and a * before or after
!      implicit double precision (a-h,o-z)
!      implicit none
!      integer ip,mult,npow
!      character string*(*),tps*(*)

```

8.19.9 Enter a TP function interactively

```

subroutine enter_tpfun_interactivly(cline,ip,longline,jp)
! interactive input of a TP expression, whole function returned in longline
!      implicit double precision (a-h,o-z)
!      implicit none
!      integer ip,jp
!      character cline*(*),longline*(*)

```

8.19.10 Deallocate TP function

Deallocates a TP function with result arrays in all equilibria.

```

subroutine tpfun_deallocate
! deallocates all arrays associated with a TP function

```

8.19.11 Enter a dummy TP function

```

subroutine enter_tpfun_dummy(symbol)
! creates a dummy entry for a TP function called symbol, used when entering
! TPfuns from a TDB file where they are not in order
!      implicit none
!      character*(*) symbol

```

8.19.12 Enter a TP function

```

subroutine enter_tpfun(symbol,text,lrot,fromtdb)
! creates a data structure for a TP function called symbol with several ranges
! text is whole expression

```

```

! lrot is returned as index.  If fromtdb is FALSE and lrot<0 it is a new
!                               expression for an old symbol
! if fromtdb is TRUE references to unknown functions are allowed
! default low temperature limit is 298.16; high 6000
  implicit none
  integer lrot
  character*(*) text,symbol
  logical fromtdb

```

8.19.13 Some more utilities for TP function

One handles nested TP function, other abbreviated names.

```

subroutine nested_tpfun(lrot,tpval,nyrot)
! called from ct1efn when a it calls another TP function that must be
! evaluated.  nyrot is the link to the ct1efn in the correct range
!  implicit double precision (a-h,o-z)
  implicit none
  integer lrot
  double precision tpval(2)
  TYPE(tpfun_expression), pointer :: nyrot
! use lowest range for all T values lower than first upper limit
! and highest range for all T values higher than the next highest limit
! one should signal if T is lower than lowest limit or higher than highest
! used  saved results if same T and P
-----
logical function compare_abbrev(name1,name2)
! returns TRUE if name1 is an abbreviation of name2
! termintaes when a space is found in name1
! each part between _ or - can be abbreviated from the left
! case insensitive. Only 36 first characters compared
  implicit none
  character*(*) name1,name2

```

8.19.14 Routines for optimizing coefficients and some other things

They are utility routines for the assessment program to manipulate the coefficients in the models.

```

subroutine enter_optvars(firstindex)
! enter variables for optimization A00-A99
  implicit none
  integer firstindex
-----
subroutine find_tpsymbol(name,type,value)

```

```

! enter variables
  implicit none
! type=0 if function, 1 if variable, 2 if optimizing variable
  integer type
  character name*(lenfnsym)
  double precision value
-----
  subroutine enter_tpconstant(symbol,value)
! enter variables
  implicit none
  character symbol*(lenfnsym)
  double precision value
-----
  subroutine change_optcoeff(lrot,value)
! change value of optimizing coefficient.  lrot is index
! -1 means just force recalculate
  implicit none
  integer lrot
  double precision value
-----
  subroutine force_recalculate_tpfuns
! force recalculation of all tpfuns by incrementing an integer in tpfuns
-----
  subroutine get_value_of_constant_name(symbol,lrot,value)
! get value (and index) of a TP constant.  lrot is index
  implicit none
  integer lrot
  character symbol*(*)
  double precision value
-----
  subroutine get_value_of_constant_index(lrot,value)
! get value of a TP constant at known lrot
  implicit none
  integer lrot
  double precision value
-----
  subroutine get_all_opt_coeff(values)
! get values of all optimizing coefficients
  implicit none
  double precision values(*)
-----
  subroutine delete_all_tpfuns
! delete all TPFUNs.  No error if some are already deleted ...
! note: tpres is deallocated when deleting equilibrium record

```

8.19.15 Saving and reading unformatted TP funs

These are used to save and read TP functions from an unformatted file.

```
subroutine save0tpfun(lfun,iws,jfun)
! save one tpfun (or parameter) with index jfun in workspace iws
!   implicit double precision (a-h,o-z)
  implicit none
  integer lfun,iws(*),jfun
-----
subroutine read0tpfun(lfun,iws,jfun)
! read one TPfun from workspace
  implicit none
  integer lfun,jfun,iws(*)
```

8.19.16 Routines used during assessments and debugging

Used to list functions using a coefficient and some other things.

```
subroutine makeoptvname(name,indx)
  implicit none
  character name*(*)
  integer indx
-----
subroutine findtpused(lfun,string)
! this routine finds which other TPFUNS (including parameters) that
! use the TPFUN lfun. It is used when listing optimizing coefficients
  implicit none
  integer lfun
  character string*(*)
-----
subroutine list_tpfun_details(lfun)
! listing the internal datastructure of all tpfuns
! converts all TP functions to arrays of coefficients with powers of T
  implicit none
  integer lfun
```

8.19.17 Routines used writing SOLGASMIX files

In FactSage and various software based on SOLGASMIX it is not possible to enter functions which can call each other and be used as model parameters. The routines below were added to write a DAT file with only coefficients.

```
subroutine tpfun2coef(ctpf,ntpf,npows,text)
```

```

! called by saveadatformat in gtp3C to generate SOLGASMIX DAT files
! converts all TP functions to arrays of coefficients with powers of T
  implicit none
  integer ntpf,npows
  type(gtp_tpfun2dat) :: ctpf(*)
  character text*(*)
-----
  subroutine list_tpascoef(lut,text,paratyp,i1,npows,factor,ctpf)
! writes a parameter in DAT format
! text contains the stoichiometries written with the format 1x,F11.6
! it can be very long if there are many coefficients.
  implicit none
  integer lut,i1,npows,paratyp
  character text*(*)
! this is a factor that may be multiplied with all coefficients for
! phases like sigma which has only a disordered part. Also ionic liquid
  double precision factor
  type(gtp_tpfun2dat) :: ctpf(*)
-----
  subroutine tpf2c(ctpf,lfun,done)
! convert TPfun lfun to an array of coefficients with powers of T
! if this TP function already converted just return
! if this TP function calls another TP function not converted return error
!
  implicit none
  integer lfun
  logical done
  type(gtp_tpfun2dat) :: ctpf(*)
-----
  subroutine tpf2cx(ctpf,lfun,nrange,cfun1)
! convert TPfun lfun to an array of coefficients with powers of T
! if this TP function already converted just return
! if this TP function calls another TP function not converted return error
  implicit none
  integer lfun,nrange
  type(gtp_tpfun_as_coeff) :: cfun1
  type(gtp_tpfun2dat) :: ctpf(*)
!   type(gtp_equilibrium_data), pointer :: ceq
-----
  subroutine adjust1range(lfun,nr1,nrange,krange,ctp1,nr2,ctp2)
! check if ctp1 range nr1 must be split in more ranges due to tbreaks in ctp2
! nrange is the total number of ranges of ctp1
! There are 10 ranges allocated for all, nr1 and nr2 are the used ranges
  implicit none
  integer lfun,nr1,nr2,krange,nrange
  type(gtp_tpfun_as_coeff) :: ctp1,ctp2

```

```

-----
subroutine adjustranges(nr1,ctp1,nr2,ctp2)
! add ctp2 to ctp1 which to have the same ranges and breakpoints
! nr1 and nr2 give the number of T-ranges and breakpoints
! add coefficients of ctp1 and ctp2 for each range
! NOTE these already multiplied with the coefficients!!
! There are 10 coefficients allocated for all functions.
! the added function is returned as ctp1
  implicit none
  integer nr1,nr2
  type(gtp_tpfun_as_coeff) :: ctp1,ctp2
-----

subroutine add1tpcoeffs(i1,ctp1,i2,ctp2,i3,ctp3)
! ctp3 is created with added coefficients from range i1 in ctp1
! and range i2 in ctp2 with same tpower. Normally i3=1
  implicit none
  integer i1,i2,i3
  type(gtp_tpfun_as_coeff) :: ctp1,ctp2,ctp3
-----

subroutine checkpowers(nc1,lfun,tpow1,npow,usedpow)
! check powers used in TP functions
! There can be several terms with same power ...
! nc1 is the maximal number of coefficients for each range (maxnc in fact)
  implicit none
  integer tpow1(*),nc1,lfun,usedpow(*),npow
-----

subroutine sortcoeffs(nc1,lfun,coeff1,tpow1)
! sort the coefficients in order power: 0 1 TlnT 2 3 -1; 7 -9 -2 other1 other2
!
!           1 2 3   4 5 6; 7 8 9   10   11
! other powers are 3, 4, -8 (meaning sqrt(t)) and maybe more
! tpow1 is array giving the T power for coeffi, 101 means T*ln(T)
! ANY CHANGE OF POWERS MUST BE MADE ALSO IN ... CHECKPOWERS
! There can be several terms with same power ...
! nc1 is the maximal number of coefficients for each range (maxnc in fact)
  implicit none
  integer tpow1(*),nc1,lfun
  double precision coeff1(*)

```

9 Iso-C declaration of OC data structures for C++

This is the file OC-isoC.h that can be used in C++ program to access data structures inside the OC software.

I would like to have some explanaton of this because I do not understand C++.

```
#if !defined __OCASI__
#define __OCASI__
```

```
/* Modification history
160829 Bo Sundman Update
2015-2016 Matthias Stratmann and Cristophe Sigli Modifications
2014 Teslos? First version
```

This contains the structure of TYPE variables in OC needed for the OC/TQ OCASI interface

NOTE there is also a c_gtp_equilibrium_data structure defined in liboctqisoc.F90 */

```
typedef struct {
    int forcenewcalc;
    double tpused[2];
    double results[6];
} tpfun_parres;
```

```
typedef struct {
    int splink, phlink, status;
    char refstate[16];
    int *endmember;
    double tpref[2];
    double chempot[2];
    double mass, molat;
} gtp_components;
```

```
typedef struct {
    int lokph, compset, ixphase, lokvares, nextcs;
} gtp_phasetuple;
```

```
typedef struct {
    int statevarid, norm, unit, phref, argtyp;
    int phase, compset, component, constituent;
    double coeff;
    int oldstv;
} gtp_state_variable;
```

```
typedef struct {
    int latd, ndd, tnoofxfr, tnoofyfr, varreslink, totdis;
    char id;
    double *dsites;
    int *nooffr;
    int *splink;
    int *y2x;
    double *dxidyj;
```



```

    double fsites;
} gtp_fraction_set;

//struct gtp_fraction_set;

typedef struct {
    int nextfree, phlink, status2, phstate, phtupx;
    double abnorm[3];
    char prefix[4], suffix[4];
    int *constat;
    double *yfr;
    double *mmyfr;
    double *sites;
    double *dpqdy;
    double *d2pqdvay;
    //struct gtp_fraction_set disfra;
    double amfu, netcharge, dgm;
    int nprop;
    int *listprop;
    double **gval;
    double ***dgval;
    double **d2gval;
    double curlat[3][3];
    double **cinvy;
    double *cxmol;
    double **cdxmol;
    double *addg;
} gtp_phase_varres;

typedef struct gtp_condition {
    int noofters, statev, active, iunit, nid, iref, seqz, experimenttype;
    int symlink1, symlink2;
    int **indices;
    double *condcoeff;
    double *prescribed, current, uncertainty;
    // should this be a struct ??
    gtp_state_variable *statvar;
    struct gtp_condition *next, *previous;
} gtp_condition;

typedef struct {
    int status, multiuse, eqno, next;
    char eqname[24], comment[72];
    double tpval[2], rtn;
    double weight;
    double *svfunres;

```

```

gtp_condition *lastcondition, *lastexperiment;
gtp_components *complist;
double **compstoi, **invcompstoi;
gtp_phase_varres *phase_varres;
tpfun_parres *eq_tpres;
double *cmuval;
double xconv;
double gmindif;
int maxiter;
char eqextra[80];
int sysmatdim, nfixmu, nfixph;
int *fixmu;
int *fixph;
double **savesysmat;
} gtp_equilibrium_data;

#endif

```

10 Tables with all 383 functions and subroutines

Name	File
double precision function evaluate_svfun_old	gtp3F.F90
double precision function mass_of	gtp3A.F90
integer function ceqrecsize	gtp3E.F90
integer function ceqsize	gtp3Y.F90
integer function degrees_of_freedom	gtp3C.F90
integer function find_phasetuple_by_indices	gtp3A.F90
integer function get_mpi_index	gtp3Y.F90
integer function get_phase_status	gtp3G.F90
integer function get_phtuplearray	gtp3A.F90
integer function getmqindex	gtp3Y.F90
integer function ispdbkeyword	gtp3E.F90
integer function istdbkeyword	gtp3E.F90
integer function newhighcs	gtp3B.F90
integer function noconst	gtp3A.F90
integer function noel	gtp3A.F90
integer function noeq	gtp3A.F90
integer function noofcs	gtp3A.F90
integer function noofphasetuples	gtp3A.F90
integer function nooftup	gtp3A.F90
integer function noph	gtp3A.F90
integer function nosp	gtp3A.F90
integer function nosusphcs	gtp3A.F90
integer function nosvf	gtp3A.F90
integer function notpf	gtp3Z.F90
integer function phvarlok	gtp3Y.F90
integer function test_phase_status	gtp3G.F90
integer function vssize	gtp3Y.F90
logical function allotropes	gtp3Y.F90
logical function allowenter	gtp3Y.F90
logical function check_minimal_ford	gtp3B.F90
logical function compare_abbrev	gtp3Z.F90
logical function fixedcomposition	gtp3Y.F90
logical function global_equil_check1	gtp3Y.F90
logical function gtp_error_message	gtp3C.F90
logical function iskeyword	gtp3E.F90
logical function ocv	gtp3Y.F90
logical function proper_symbol_name	gtp3Y.F90
logical function same_statevariable	gtp3D.F90
logical function same_stoik	gtp3Y.F90
logical function test_phase_status_bit	gtp3G.F90
logical function testelstat	gtp3G.F90

Name	File
logical function testspstat	gtp3G.F90
subroutine add1tpcoeffs	gtp3Z.F90
subroutine add_addrrecord	gtp3H.F90
subroutine add_fraction_set	gtp3G.F90
subroutine add_to_reference_phase	gtp3G.F90
subroutine addition_selector	gtp3H.F90
subroutine adjustlrange	gtp3Z.F90
subroutine adjustranges	gtp3Z.F90
subroutine alphaelorder	gtp3G.F90
subroutine alphaphorder	gtp3G.F90
subroutine alphaspororder	gtp3G.F90
subroutine amend_components	gtp3A.F90
subroutine amend_global_data	gtp3D.F90
subroutine any_disordered_part	gtp3E.F90
subroutine apply_condition_value	gtp3D.F90
subroutine ask_default_constitution	gtp3D.F90
subroutine ask_phase_constitution	gtp3D.F90
subroutine ask_phase_new_constitution	gtp3D.F90
subroutine assessmenthead	gtp3A.F90
subroutine bccendmem	gtp3B.F90
subroutine bccint1	gtp3B.F90
subroutine bccint2	gtp3B.F90
subroutine bccpermut	gtp3B.F90
subroutine biblioread	gtp3E.F90
subroutine bibliosave	gtp3E.F90
subroutine calc_debyecp	gtp3H.F90
subroutine calc_diffusion	gtp3H.F90
subroutine calc_disfrac	gtp3X.F90
subroutine calc_disfrac2	gtp3X.F90
subroutine calc_einsteincp	gtp3H.F90
subroutine calc_elastica	gtp3H.F90
subroutine calc_gridpoint	gtp3Y.F90
subroutine calc_magnetic_inden	gtp3H.F90
subroutine calc_molmass	gtp3F.F90
subroutine calc_phase_mol	gtp3F.F90
subroutine calc_phase_molmass	gtp3F.F90
subroutine calc_qf	gtp3F.F90
subroutine calc_schottky_anomaly	gtp3H.F90
subroutine calc_secondeinstein	gtp3H.F90
subroutine calc_twostate_model1	gtp3H.F90
subroutine calc_twostate_model_john	gtp3H.F90

Name	File
subroutine calc_twostate_model_old	gtp3H.F90
subroutine calc_volmod1	gtp3H.F90
subroutine calc_xiongmagnetic	gtp3H.F90
subroutine calcg	gtp3X.F90
subroutine calcg_endmember	gtp3Y.F90
subroutine calcg_endmember6	gtp3Y.F90
subroutine calcg_endmemberx	gtp3Y.F90
subroutine calcg_internal	gtp3X.F90
subroutine calculate_reference_state	gtp3F.F90
subroutine calculate_reference_state_old	gtp3F.F90
subroutine cgint	gtp3X.F90
subroutine change_element_status	gtp3G.F90
subroutine change_many_phase_status	gtp3G.F90
subroutine change_optcoeff	gtp3Z.F90
subroutine change_phase_status	gtp3G.F90
subroutine change_phtup_status	gtp3G.F90
subroutine change_species_status	gtp3G.F90
subroutine check_alphaindex	gtp3G.F90
subroutine checkdb	gtp3E.F90
subroutine checkdb2	gtp3E.F90
subroutine checkdefcon	gtp3Y.F90
subroutine checkpowers	gtp3Z.F90
subroutine clear_phase_status_bit	gtp3G.F90
subroutine compmassbug	gtp3Y.F90
subroutine condition_value	gtp3D.F90
subroutine config_entropy	gtp3X.F90
subroutine config_entropy_cqc6	gtp3X.F90
subroutine config_entropy_cqc_classicqc	gtp3X.F90
subroutine config_entropy_floryhuggins	gtp3X.F90
subroutine config_entropy_i2sl	gtp3X.F90
subroutine copy_condition	gtp3B.F90
subroutine copy_equilibrium	gtp3B.F90
subroutine copy_equilibrium2	gtp3B.F90
subroutine copy_fracset_record	gtp3G.F90
subroutine create_constitlist	gtp3G.F90
subroutine create_debyecp	gtp3H.F90
subroutine create_diffusion	gtp3H.F90
subroutine create_einsteincp	gtp3H.F90
subroutine create_elastic_model_a	gtp3H.F90
subroutine create_endmember	gtp3G.F90
subroutine create_interaction	gtp3G.F90

Name	File
subroutine create_magrec_inden	gtp3H.F90
subroutine create_parrecords	gtp3G.F90
subroutine create_proprec	gtp3G.F90
subroutine create_schottky_anomaly	gtp3H.F90
subroutine create_secondeinstein	gtp3H.F90
subroutine create_twostate_model1	gtp3H.F90
subroutine create_volmod1	gtp3H.F90
subroutine create_xiong_magnetic	gtp3H.F90
subroutine ct1efn	gtp3Z.F90
subroutine ct1getsym	gtp3Z.F90
subroutine ct1mexpr	gtp3Z.F90
subroutine ct1mfn	gtp3Z.F90
subroutine ct1power	gtp3Z.F90
subroutine ct1wfn	gtp3Z.F90
subroutine ct1wpow	gtp3Z.F90
subroutine ct1xfn	gtp3Z.F90
subroutine ct2mfn	gtp3Z.F90
subroutine deallocate_gtp	gtp3E.F90
subroutine decode_constarr	gtp3C.F90
subroutine decode_state_variable	gtp3F.F90
subroutine decode_state_variable3	gtp3F.F90
subroutine decode_stoik	gtp3C.F90
subroutine delete_all_conditions	gtp3B.F90
subroutine delete_all_tpfuns	gtp3Z.F90
subroutine delete_equilibria	gtp3B.F90
subroutine delphase	gtp3E.F90
subroutine diffusion_onoff	gtp3H.F90
subroutine disordery	gtp3X.F90
subroutine disordery2	gtp3X.F90
subroutine elements2components1	gtp3G.F90
subroutine encode_constarr	gtp3C.F90
subroutine encode_state_variable	gtp3F.F90
subroutine encode_state_variable3	gtp3F.F90
subroutine encode_state_variable_record	gtp3F.F90
subroutine encode_stoik	gtp3C.F90
subroutine enter_bibliography_interactively	gtp3D.F90
subroutine enter_composition_set	gtp3B.F90
subroutine enter_default_constitution	gtp3D.F90
subroutine enter_element	gtp3B.F90
subroutine enter_equilibrium	gtp3B.F90
subroutine enter_experiment	gtp3D.F90

Name	File
subroutine enter_many_equil	gtp3B.F90
subroutine enter_material	gtp3D.F90
subroutine enter_optvars	gtp3Z.F90
subroutine enter_parameter	gtp3B.F90
subroutine enter_parameter_interactively	gtp3D.F90
subroutine enter_phase	gtp3B.F90
subroutine enter_species	gtp3B.F90
subroutine enter_species_property	gtp3D.F90
subroutine enter_svfun	gtp3F.F90
subroutine enter_tpconstant	gtp3Z.F90
subroutine enter_tpfun	gtp3Z.F90
subroutine enter_tpfun_dummy	gtp3Z.F90
subroutine enter_tpfun_interactively	gtp3Z.F90
subroutine enterphase	gtp3B.F90
subroutine eval_tpfun	gtp3Z.F90
subroutine evaluate_all_svfun_old	gtp3F.F90
subroutine expand_wildcards	gtp3E.F90
subroutine extend_proprec	gtp3G.F90
subroutine extract_massbalcond	gtp3X.F90
subroutine extract_stvr_of_condition	gtp3D.F90
subroutine fccint211	gtp3B.F90
subroutine fccint22	gtp3B.F90
subroutine fccint31	gtp3B.F90
subroutine fccip2A	gtp3B.F90
subroutine fccip2B	gtp3B.F90
subroutine fccpe1111	gtp3B.F90
subroutine fccpe211	gtp3B.F90
subroutine fccpermut	gtp3B.F90
subroutine find_component_by_name	gtp3A.F90
subroutine find_constituent	gtp3A.F90
subroutine find_defined_property	gtp3C.F90
subroutine find_element_by_name	gtp3A.F90
subroutine find_gridmin	gtp3Y.F90
subroutine find_phase_by_name	gtp3A.F90
subroutine find_phase_by_name_exact	gtp3A.F90
subroutine find_phasetuple_by_name	gtp3A.F90
subroutine find_phasex_by_name	gtp3A.F90
subroutine find_species_by_name	gtp3A.F90
subroutine find_species_record	gtp3A.F90
subroutine find_species_record_exact	gtp3A.F90
subroutine find_species_record_noabbr	gtp3A.F90

Name	File
subroutine find_svfun	gtp3F.F90
subroutine find_symbol_with_equilno	gtp3F.F90
subroutine find_tpfun_by_name	gtp3Z.F90
subroutine find_tpfun_by_name_exact	gtp3Z.F90
subroutine find_tpsymbol	gtp3Z.F90
subroutine findconst	gtp3B.F90
subroutine findeq	gtp3A.F90
subroutine findtpused	gtp3Z.F90
subroutine force_recalculate_tpfuns	gtp3Z.F90
subroutine format_phase_composition	gtp3C.F90
subroutine geneqname	gtp3B.F90
subroutine generate_charged_grid	gtp3Y.F90
subroutine generate_dense_grid	gtp3Y.F90
subroutine generate_fccord_grid	gtp3Y.F90
subroutine generate_grid	gtp3Y.F90
subroutine generate_ionliq_grid	gtp3Y.F90
subroutine generic_grid_generator	gtp3Y.F90
subroutine get_all_conditions	gtp3C.F90
subroutine get_all_opt_coeff	gtp3Z.F90
subroutine get_component_name	gtp3A.F90
subroutine get_condition	gtp3D.F90
subroutine get_condition2	gtp3D.F90
subroutine get_constituent_location	gtp3A.F90
subroutine get_constituent_name	gtp3A.F90
subroutine get_diffusion_matrix	gtp3H.F90
subroutine get_element_data	gtp3A.F90
subroutine get_experiment_with_symbol	gtp3D.F90
subroutine get_many_svar	gtp3F.F90
subroutine get_one_condition	gtp3C.F90
subroutine get_one_experiment	gtp3C.F90
subroutine get_parameter_typty	gtp3D.F90
subroutine get_phase_compset	gtp3A.F90
subroutine get_phase_data	gtp3A.F90
subroutine get_phase_name	gtp3A.F90
subroutine get_phase_record	gtp3A.F90
subroutine get_phase_structure	gtp3A.F90
subroutine get_phase_variance	gtp3A.F90
subroutine get_phasetup_name	gtp3A.F90
subroutine get_phasetup_name_old	gtp3A.F90
subroutine get_phasetup_record	gtp3A.F90
subroutine get_species_component_data	gtp3A.F90

Name	File
subroutine get_species_data	gtp3A.F90
subroutine get_species_location	gtp3A.F90
subroutine get_species_name	gtp3A.F90
subroutine get_stable_state_var_value	gtp3F.F90
subroutine get_state_var_value	gtp3F.F90
subroutine get_value_of_constant_index	gtp3Z.F90
subroutine get_value_of_constant_name	gtp3Z.F90
subroutine global_gridmin	gtp3Y.F90
subroutine gtpread	gtp3E.F90
subroutine gtpsave	gtp3E.F90
subroutine gtpsavedir	gtp3E.F90
subroutine gtpsavelatex	gtp3E.F90
subroutine gtpsavetdb	gtp3E.F90
subroutine gtpsavetm	gtp3E.F90
subroutine gtpsaveu	gtp3E.F90
subroutine incunique	gtp3E.F90
subroutine init_gtp	gtp3A.F90
subroutine intsort	gtp3E.F90
subroutine line_with_phases_withdgm0	gtp3C.F90
subroutine list_addition	gtp3H.F90
subroutine list_addition_values	gtp3H.F90
subroutine list_all_components	gtp3C.F90
subroutine list_all_elements	gtp3C.F90
subroutine list_all_elements2(unit)	gtp3C.F90
subroutine list_all_funs	gtp3Z.F90
subroutine list_all_phases	gtp3C.F90
subroutine list_all_species	gtp3C.F90
subroutine list_all_svfun	gtp3F.F90
subroutine list_bibliography	gtp3C.F90
subroutine list_components_result	gtp3C.F90
subroutine list_conditions	gtp3C.F90
subroutine list_defined_properties	gtp3C.F90
subroutine list_element_data	gtp3C.F90
subroutine list_equilibria_details	gtp3C.F90
subroutine list_free_lists	gtp3Y.F90
subroutine list_global_results	gtp3C.F90
subroutine list_many_formats	gtp3C.F90
subroutine list_phase_data	gtp3C.F90
subroutine list_phase_data2	gtp3C.F90
subroutine list_phase_model	gtp3C.F90
subroutine list_phase_results	gtp3C.F90

Name	File
subroutine list_phases_with_positive_dgm	gtp3C.F90
subroutine list_short_results	gtp3C.F90
subroutine list_sorted_phases	gtp3C.F90
subroutine list_species_data	gtp3C.F90
subroutine list_species_data2	gtp3C.F90
subroutine list_svfun	gtp3F.F90
subroutine list_tpascoef	gtp3Z.F90
subroutine list_tpfun	gtp3Z.F90
subroutine list_tpfun_details	gtp3Z.F90
subroutine list_unentered_funs	gtp3Z.F90
subroutine listoptcoef	gtp3C.F90
subroutine locate_condition	gtp3D.F90
subroutine lower_case_species_name	gtp3E.F90
subroutine make_stvrec	gtp3F.F90
subroutine makeoptvname	gtp3Z.F90
subroutine mark_stable_phase	gtp3G.F90
subroutine merge_gridpoints	gtp3Y.F90
subroutine need_propertyid	gtp3H.F90
subroutine nested_tpfun	gtp3Z.F90
subroutine new_gtp	gtp3E.F90
subroutine palmtree	gtp3Y.F90
subroutine pop_pyval	gtp3X.F90
subroutine push_pyval	gtp3X.F90
subroutine read0tpfun	gtp3Z.F90
subroutine readash	gtp3E.F90
subroutine readendmem	gtp3E.F90
subroutine readequil	gtp3E.F90
subroutine readintrec	gtp3E.F90
subroutine readpdb	gtp3E.F90
subroutine readphases	gtp3E.F90
subroutine readproprec	gtp3E.F90
subroutine readtdb	gtp3E.F90
subroutine readtdbsilent	gtp3E.F90
subroutine remove_composition_set	gtp3B.F90
subroutine replacetab	gtp3E.F90
subroutine restore_constitutions	gtp3X.F90
subroutine restore_phases_implicitly_suspended	gtp3G.F90
subroutine restore_species_implicitly_suspended	gtp3G.F90
subroutine save0tpfun	gtp3Z.F90
subroutine save_constitutions	gtp3X.F90
subroutine save_datformat	gtp3E.F90

Name	File
subroutine save_results	gtp3G.F90
subroutine saveash	gtp3E.F90
subroutine saveequil	gtp3E.F90
subroutine savephases	gtp3E.F90
subroutine select_composition_set	gtp3Y.F90
subroutine selecteq	gtp3A.F90
subroutine separate_constitutions	gtp3Y.F90
subroutine set_cond_or_exp	gtp3D.F90
subroutine set_condition	gtp3D.F90
subroutine set_constituent_reference_state	gtp3G.F90
subroutine set_constitution	gtp3A.F90
subroutine set_default_constitution	gtp3Y.F90
subroutine set_driving_force	gtp3X.F90
subroutine set_emergency_startpoint	gtp3Y.F90
subroutine set_input_amounts	gtp3D.F90
subroutine set_lattice_parameters	gtp3H.F90
subroutine set_metastable_constitutions2	gtp3Y.F90
subroutine set_new_stoichiometry	gtp3A.F90
subroutine set_phase_amounts	gtp3Y.F90
subroutine set_phase_status_bit	gtp3G.F90
subroutine set_putfun_constant	gtp3F.F90
subroutine set_reference_state	gtp3A.F90
subroutine set_uniquac_species	gtp3D.F90
subroutine set_unit	gtp3G.F90
subroutine setendmemarr	gtp3X.F90
subroutine setpermolebit	gtp3H.F90
subroutine shiftcompsets	gtp3Y.F90
subroutine sort_ionliqconst	gtp3B.F90
subroutine sortcoeffs	gtp3Z.F90
subroutine sortinphtup	gtp3Y.F90
subroutine state_variable_val	gtp3F.F90
subroutine state_variable_val3	gtp3F.F90
subroutine store_putfun	gtp3F.F90
subroutine store_putfun_old	gtp3F.F90
subroutine subrefstates	gtp3C.F90
subroutine sumprops	gtp3F.F90
subroutine suspend_composition_set	gtp3B.F90
subroutine suspend_phases_implicitly	gtp3G.F90
subroutine suspend_species_implicitly	gtp3G.F90
subroutine svfunread	gtp3E.F90
subroutine svfunsave	gtp3E.F90

Name	File
subroutine switch_compsets2	gtp3Y.F90
subroutine tabder	gtp3X.F90
subroutine tdbrefs	gtp3B.F90
subroutine termterm	gtp3G.F90
subroutine todo_after_found_equilibrium	gtp3Y.F90
subroutine todo_before	gtp3Y.F90
subroutine tpf2c	gtp3Z.F90
subroutine tpf2cx	gtp3Z.F90
subroutine tpfun2coef	gtp3Z.F90
subroutine tpfun_deallocate	gtp3Z.F90
subroutine tpfun_init	gtp3Z.F90
subroutine uniquac_model	gtp3X.F90
subroutine verify_phase_varres_array	gtp3Y.F90
subroutine write_pdbformat	gtp3E.F90

11 Summary

Thats all! Be happy YOU did not have to write all this.

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