# User Guide to the OpenCalphad software package version 5.0

DRAFT Bo Sundman, August 21, 2018



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# 1 Introduction

The development of the OpenCalphad (OC) sofware was started by a small group of dedicated scientists who wanted to provide an open source multicomponent thermodynamic software. It aims to provide a free high quality software for thermodynamic calculations, including property and phase diagrams, assessment of databases and a thermodynamic library for simulations for inorganic systems i.e. gases. liquids, alloys and other materials using many different kinds of models for the phases. There are three basic papers published about OC [15Sun1, 15Sun2, 16Sun]. General information about thermodynamic models, calculations and assessments based on the Calphad technique can be found in the book by Lukas et al [07Luk]. The software is provided free with a GNU GPL license.

In OC there is also a framework to store different kinds of materials properties that depend on temperature, pressure and composition when such properties are related to the phases of the system. The OC software can also be used to assess model parameters for such properties from experimental and theoretical values.

Complimentary (and maybe sometimes contradictory, I am not perfect) information about the OC software can be found in getting-started.pdf, news-oc5.pdf and the other parts of the OC documentation.

# 2 Some general features

The different parts of the OC software are documented separately for each module: thermodynamic models (GTP), equilibrium calculations (HMS), step/map/plot routines (SMP) and the application software interface (OCASI/TQ). OC uses the free numerics packages LAPACK and BLAS and a free least square routine, LMDIF [LMDIF], developed at Argonne 1980, for assessments. For graphics OC generates a command file which can be plotted with the free GNUPLOT [GNUPLOT] software. If GNUPLOT is properly installed this is done automatically by OC.

### 2.1 Command line user interface

OC is operated by commands typed by the user or read from a macro file. The command monitor has a menu of command and each of these usually has sub-menus and finally some questions may be asked like phase names, a value or an expression. In most cases a default answer is provided which can be selected by just pressing the RETURN key or by typing a comma, ",", on the same line as the command. At all levels the user should be able to type a ? and get some help, usually an extract from this manual, sometimes just a menu or examples of answers.

If you prefer a graphical user interface (GUI) you are welcome to add this to the OC software.

### 2.1.1 Popup window for read/save

There is a problem using a command line interface when you want to open a file for reading or saving unless this file is on the same directory as where you started the program. Thus from version 5.018 there is a popup window to open a file (for a macro, a database or to save a calculation). In this window you can browse your directories to find the file.

This has some consequencies for editing your macro files which you should be aware of and which are explained below. The code for the popup window, tinyfiledialogs, is written in C by Guillaume Vareille.

You can turn off the open file popup window feature with the command set advanced popup Y. You can turn it on again with the same command finishing with anything but Y.

- The directory where you start the session with OC is called the "working directory". On a linux system you can find this directory by typing "pwd" before starting OC (or if you type @pwd inside OC). On a Windows system you can see the working directory and its files if you type @dir inside OC.
- When the popup window is opened the directories and files matching the "filter" in the working directory should be listed. If not you can select a directory inside the popup window. The filter when open a macro file is "OCM" and when opening a database file it is "TDB" which means only files with these extensions are listed. You can change the directory in the popup window to select the file you want and you can read a file with another extension. OC will save internally the directory where you start the macro.
- Inside a macro file you normally read a TDB file and if you do not specify the name of the database on the same line as the command *read tdb* the popup window will open so you can specify the file in this window.
- But normally you know which database you want to use inside the macro and if you give the file name on the same line as the commad: read tdb filename the popup window will not open and OC will search for the specified database file starting from the "working directory". But if the database file is in the same directory as the macro file you MUST prefix "filename" with "./", i.e. read tdb ./filename. You may include directories in "filename", (including "../" to go to the directory above). OC will replace the "./" by the directory where you started the macro or prefix "../" by this directory.
- In the macro file you can give the full path to the file to be opened but that is rather clumsy.
- When you open a file for write inside a macro, like output from a plot, you can also specify the file name in the command prefixed by "./" if you want to save the file on the same directory as the macro file. Otherwise it will be saved at the working directory.
- If you use the switch "/output=" or "/append=" after a command to redirect output from the command you can also use the popup window to specify the file name or use a filename with or without the prefix "./". The filter in this case is "DAT".

Opening files on different directories can be complicated inside OC. For example during assessments you may use many different files for generating graphics and unformatted save files. Preferably you keep all of these on the same directory.

You are welcome to provide feedback on this popup feature and other parts of the user interface.

# 2.2 Names and symbols

There are many symbols and names used in this package. A symbol or name MUST start with a letter A-Z. It usually can contain digits and the underscore character after the initial letter. All names are CASE INSENSITIVE, i.e. fe, FE, fE and Fe is the same. Some special symbols are used:

- /- is used to denote the electron. /+ or /- -1 can be used for a positive charge.
- $\bullet$  # are used to identify composition sets after a phase name or sublattice after a constituent name.
- & are used in some parameter identifiers to specify the constituent for the parameter, like for mobilities, the mobility of Fe in the BCC phase is denoted MQ&FE(BCC).

A name of an element is one or two characters, a species maximum 24 characters (note that a species name does not have to be its stoichiometric formula). A phase name is 24 characters but can also have a pre- and suffix 4 characters long and possibly a composition set number after a hash symbol, #.

State variable symbols and TP-fun symbols can be 16 characters long. TP-funs are expressions used to describe the T and P dependence of model parameters.

For user input it is possible to use abbreviations of names but you must be careful with names that have the same abbreviation and avoid phase names that are abbreviations of another phase!

# 2.3 Phases, composition sets and phase tuples

Each phase in a system has a name and a thermodynamic model. The models are explained in a separate documentation. The phases can be entered interactively or read from a database or a saved file together with the last calculation.

In some cases a phase can be stable with two ore more different compositions for example inside miscibility gaps or when the phase has order/disorder transitions. In such a case you use a composition set index to separate these. The composition set index is appended to the phase name preceded by a hash "#" character, like liquid#2.

Composition sets can be created manually, see the command **AMEND PHASE** in section 5.10.4 or automatically by the grid minimizer or application software.

The phase tuple has been introduced to have a single index for both phases and composition sets in application software. The tuple index thus contain both the phase number and the composition set index. The array of tuple indices is updated internally whenever a new composition set is created or deleted.

### 2.4 The use of wildcards for names

In many cases you can use an asterix "\*" as a name and this normally means "all". For setting status of phases you can use the special "\*S" for all suspended phase, "\*D" for all dormant phases.

### 2.5 State variables

A state variable in a thermodynamic system has a value which at equilibrium is independent of the way the system has reach its current state. All state variables available in OC are listed in Table 1. They are used to set conditions and to obtain results from an equilibrium calculation. It is possible to use state variables also when close to the equilibrium state for example when simulating a phase transformation.

# 2.6 Model parameters

All data is organized relative to a phase and the phase is identified by a name. Each phase can have a different model for the composition dependence but the way to enter model parameters is the same for all models. However, the meaning of a model parameter will depend on the model of the phase.

Many types of data can be stored as explained in the section on parameter identifiers. The parameter also has a constituent specification explained in the constituent array section and possibly a degree, the meaning of which is model dependent and a bibliographic reference.

The basic syntax of a parameter is

"identifier" ("phase name", "constituent array"; "degree") "expression" "bibl.ref."

These parts are explained in more detail below.

### 2.6.1 Model Parameter Identifiers

The OC thermodynamic package can handle any phase property that depend on T, P and the constitution of the phase using the models implemented. It is easy to extend the number of properties by declaring property identifiers in the source code. If the parameters should have an influence on the Gibbs energy (like the Curie temperature) or a diffusion coefficient (like the mobility) the necessary code to calculate this must be added.

Table 1: A 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					
Τ	1	-	-	-	Temperature
P	2	-	-	-	Pressure
MU	3	component	-/phase	-	Chemical potential
AC	4	component	-/phase	-	Activity
LNAC	5	component	-/phase	-	LN(activity)
		Exter	nsive and norr	nallized proper	ties
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 <sup>3</sup>
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	$7\mathrm{z}$	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		rs			
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

A list of the model parameter identifiers as shown in Table 2 can be obtained by the command LIST MODEL\_PARAM\_ID

Several of these identifiers have no supporting software implemented, this is an ongoing project. The columns T P indicate if the parameter may depend on T or P. Some identifiers require additional specification of the constituent and sublattice, like the mobility of a constituent. Currently it is not yet clear if mobilities should depend on the sublattice or not

Table 2: Current set of model parameter identifiers

Indx	Ident	Т	P	Specification	Status	Note
1	G	T	P		0	Energy
2	TC	-	P		2	Combined Curie/Neel T
3	BMAG	-	-		1	Average Bohr magneton numb
4	CTA	-	P		2	Curie temperature
5	NTA	-	P		2	Neel temperature
6	IBM	-	P	& <constituent#sublattice></constituent#sublattice>	12	Individual Bohr magneton num
7	THET	-	P		2	Debye or Einstein temp
8	VO	-	-		1	Volume at TO, PO
9	VA	T	-		4	Thermal expansion
10	VB	T	P		0	Bulk modulus
11	G2	T	P		0	Liquid two state parameter
12	CBT	T	P		2	Hickel T
13	MQ	T	P	& <constituent#sublattice></constituent#sublattice>	10	LN mobility1 of component
14	MF	T	P	& <constituent#sublattice></constituent#sublattice>	10	LN mobility2 of component
15	MG	T	P	& <constituent#sublattice></constituent#sublattice>	10	LN mobility3 of component
16	THT2	T	P		2	Smooth step function T
17	DCP2	T	P		2	Smooth step function value
18	VISC	T	P		0	Viscosity
19	LPX	T	P		0	Lattice param X axis
20	LPY	T	P		0	Lattice param Y axis
21	LPZ	T	P		0	Lattice param Z axis
22	LPTH	T	P		0	Lattice angle TH
23	EC11	T	P		0	Elastic const C11
24	EC12	T	P		0	Elastic const C12
25	EC44	T	P		0	Elastic const C44
26	FHV	T	P	& <constituent#sublattice></constituent#sublattice>	10	Flory-Huggins volume ratio
27	RHO	T	P		0	Electrical resistivity
28	LAMB	T	P		0	Thermal conductivity
29	HMVA	T	P		0	Enthalpy of vacancy form
30	TSCH	-	P		2	Schottky anomality T
31	CSCH	-	P		2	Schottky anomality Cp/R

but the notation allows that.

A slightly more detailed explanation of the identifiers are:

- G, the Gibbs energy parameter for an endmember or an interaction. G(LIQUID,FE;0) is the Gibbs energy for pure liquid Fe. Note that the parameter will be used also below the melting temperature of Fe for a liquid phase containing Fe. G(LIQUID,CR,FE;0) is the regular parameter for Cr and Fe in the liquid.
- TC, a parameter for the critical temperature for ferro or antiferro magnetic ordering using the Inden model.

- BMAG, a parameter for the average Bohr magneton number using the Inden model.
- CTA, a parameter for the Curie temperature for ferromagnetic ordering using a modified Inden model.
- NTA, a parameter for the Neel temperature for antiferromagnetic ordering using a modified Inden model.
- IBM&C, a parameter for the individual Bohr magneton number for constituent C using a modified Inden model. For example IBM&FE(BCC,FE) is the Bohr magneton number for BCC Fe. The identifier IBM&FE(BCC,CR) means the Bohr magneton number of a single Fe atom in BCC Cr. An identifier IBM&FE(BCC,CR,FE) can be used to decribe the composition dependence of the Bohr magneton number for Fe in BCC.
- THET, a parameter for the Debye or Einstein temperature.
- V0, a parameter for the volume at 298.15 K and 1 bar.
- VA, a parameter for the integrated thermal expansion.
- VB, a parameter for the Bulk modulus.
- G2, a parameter for the two-state liquid model.
- CBT, a parameter for the crystal breakdown T.
- MQ&C, a parameter for the logarithm of the frequency factor of the mobility of constituent C.
- MF&C, a parameter for the activition energy of the mobility of constituent C.
- MG&C, a parameter for the magnetic factor of the mobility of constituent C.
- THT2, The T for a smooth change of  $C_P$
- DCP2, The value of the smooth change in J/mol
- VISC, a parameter for the viscosity.
- LPX, a parameter the lattice parameter in X direction.
- LPY, a parameter the lattice parameter in Y direction.
- LPZ, a parameter the lattice parameter in Z direction.
- LPTH, a parameter the angle between lattice directions.
- EC11, a parameter for the elastic constant C11.
- EC12, a parameter for the elastic constant C12.

- EC44, a parameter for the elastic constant C44.
- FHV&C, a parameter for the Flory-Huggins volume ratio of C
- RHO, a parameter for the electrical resistivity.
- LAMB, a parameter for the thermal conductivity.
- HMVA, a parameter for the enthalpy of vacancy formation.
- TSCH, the T for a Schottky anomality
- CSCH, the Schottky anomality  $\Delta C_P$ .

The current value of any of these parameter identifiers can be obtained by the command **LIST STATE\_VARIABLE** using the identifier and appropriate phase and component specifiers, see section 16.16.

For details of the meaning of the model identifier refer to the model documentation. As already mentioned many of the identifiers, like the mobility, does not influence the Gibbs energy but as they depend on the T, P and constitution of the phase it is convenient to model them in the same way as the thermodynamic data.

### 2.6.2 Constituent array and degrees

A constituent array specifies one or more constituent in each sublattice. A constituent must be entered as a species with fixed stoichiometry. Between constituents in different sublattices you must give a colon, ":", between interacting constituents in the same sublattice you must give a comma, "," or a space. A constituent array with exactly one constituent in each sublattice is also called an "endmember" as it give the value for a "compound" with fixed stoichiometry. Constituent arrays with one or more interaction constituents describe the composition dependence of the property. Without such parameters the property will vary linearly between the endmembers.

If there are no sublattices, like in the gas, you just give the phase and the constituent G(GAS,C1O2)

If no degree is specified it is assumed to be zero. For endmembers the degree must be zero but it may sometimes be useful to specify the zero in order to distinguish the parameter from the expression for the calculated value of the property, like the chemical potential of a component. In the gas phase you normally assumes there are no interactions but it is possible to add such parameters. For an fcc phase with 4 sublattice for ordering and one for interstitials an endmember parameter is

G(FCC,AL:NI:NI:NI:VA;0)

This would be the Gibbs energy of an fcc AL1NI3 ordered compound.

An interaction between vacancies and carbon in the austenite is

```
G(FCC,FE:C,VA;0)
```

For an interaction parameter you should always specify a degree but also in this case an omitted degree is interpreted as zero.

### 2.6.3 The TPFUN expression and bibliographic reference

The expression for a parameter can be a single value or a function of T and P. It must start with a low temperature limit, usually 298.15 K and must finish with a high temperature limit. These expressions as well as their first an second derivatives will be calculated by the TP-fun package. To simplify that there is a strict syntax for the expression. A term in the expression is

```
"numeric value" * "name of TP function" *T** "power" *P** "power"
```

You can construct very complex expression by referring to other functions. If "power" is zero the corresponding \*T\*\* or \*P\*\* can be omitted. If it is negative it must be surrounded by parenthesis like (-1). If it is unity the \*\*1 can be skipped.

Several terms, seperated by signs, forms an expression and it must be terminated by a semicolon, ";". After the semicolon there must be a high temperature limit or a breakpoint in temperature. A breakpoint must be followed by the letter "Y" and then a new expression for temperatures above the breakpoint.

It is the responsability of the database manager to ensure the expression is continuous at the breakpoint. If there are jumps in the value at a breakpoint strange things will happen when calculating equilibria.

After the high temperature limit the letter "N" must be given followed by a bibliographic reference for the parameter. Use the commands AMEND or ENTER BIBLIOGRAPHIC to give the reference.

The database manager should always add a bibliographic reference even if it is just his or her name and a date. This avoids people to mistake a value inspired by your experience for a carefully validated parameter.

A term can be used inside a natural logarithm, LN, or exponential, EXP. And the LN or EXP can be multiplied with a term. On the other hand you are not allowed to have any parenthesis, except around powers or arguments to LN and EXP. A valid expression is

```
298.15 -8856.94+157.48*T-26.908*T*LN(T)+.00189435*T**2
-1.47721E-06*T**3+139250*T**(-1); 2180 Y
-34869.344+344.18*T-50*T*LN(T)-2.88526E+32*T**(-9); 6000 N 91Din
```

where 91Din is the bibliographic reference to the SGTE unary database.

# 2.7 The reference state of a component

The values of most thermodynamic data must have a defined reference state. By default the reference state for the components is SER (Stable Element Reference) which is the stable state of the element at 298.15 K and 1 bar. (NOTE: the default reference state is defined by the database but today almost all databases have SER as reference state.)

For each component (also for other components than the elements) you can specify a phase at a given temperature and pressure as reference state, see section 26.17. The phase must exist for the component as pure.

A state variable like the chemical potential, MU(O), will refer to the user defined reference state if set. To obtain the value for the SER state you can use a suffix S, i.e. MUS(O) to obtain the chemical potential referred to SER. All state variables are listed in Table 1.

Note that the value of integral properties like Gibbs energy, G, enthalpy, H, etc. may have mixed reference states unless all components have the same phase as reference state. In order to have the enthalpy of mixing of a phase all components must have that phase as reference state. For the volume, V, SER is always used as reference state unless all components have the same reference state.

### 2.8 Macro files

The macro command is very useful for preparing complex calculations and to remember how you did them. A macro file is simplest to create staring from a log file (created by the **SET LOG** command). See the macros directory for examples.

After a macro command the popup window will allow you to search for the file on all your directories unless you type the name of the file on the same line. In the latter case the macro file must be on you "working directory", see section 2.1.1

When you open files, such as databases, inside a macro file and you type the file name on the same line as the command as "read tdb ./steel1", you must prefix the file name, "steel1" with "./" if the tdb file is on the same directory as the macro file. If your command line is just "read tdb" the popup window will be activated and you can specify the file there.

If you open another macro file inside a macro (typically when you do assessments) you must also prefix the name of the macro with "./" unless you want to select the macro using the popup window.

You can insert stops in the macro file with "@&" at the beginning of a line. This can be useful to inspect the output. The macro continues after pressing the ENTER/RETURN key. Depending on the graphical driver you use the program will normally pause after each plot and you must use the mouse to click on the graphical window to continue.

You can insert comments in the macro file with "@\$" at the beginning of the comment line.

A macro file should be terminated with the command **SET INTERACTIVE** which gives back control to the keyboard (or the calling macro file) otherwise the program may terminate

at the end of the macro.

Macro files can be nested 5 levels deep.

# 2.9 Assessment of model parameters

One of the important uses of the OC software is to assess model parameters in the phases of a system using experimental and theoretical data. This is done by recalculating the experimental data from the model and by varying the model parameters a least square routine, LMDIF developed at Argonne National Lab in 1981, is used to find the best set.

Assessments are a very difficult procedure as you must also take into account the extrapolations of the model outside the range of experimental data. So called "First Principles Calculations" or the somewhat simpler "Density Functional Theory" (DFT) which are based on the electronic structure of the elements can provide information for metastable as well as for the stable state. But you must be careful that the result from such calculations does not represent a mechanically unstable state with imaginary phonon frequencies.

Experimental data can be direct measurements of thermodynamic data like enthalpies, chemical potentials, heat capacities, activities, etc but very important are also measurements of phase diagrams, solubilities etc because they are also related to the equilibrium state.

There are several commands related to the assessment procedure in OC but during the assessment you will also use the basic facilities to calculate equilibria for different kinds of conditions as well as many different kinds of diagrams to verify the results.

### 2.9.1 Entering coefficients to be assessed

The command "enter optimizing coefficients", see section 10.11 creates symbols A00 up to A99 that can be used as coefficients in the thermodynamic model parameters. Maximum number of coefficients are 100.

### 2.9.2 Entering phases and model parameters

The elements, species and phases with their appropriate models are entered using the appropriate commands. Normally this is on a macro file in order to have proper documentation. Keep also in mind that an assessment is often revised after a few years when new data become available or you find that the extrapolations of an assessment to a higher order system is not reasonable.

The model parameters are entered using "enter parameter", see section 10.12 or "enter tpfun", see section 10.18 as many parameters may share some properties and a TP-function can be used in several parameters. The optimizing coefficients A00 to A99 with different T and P dependence can be used instead of numerical values as their values should be assessed.

### 2.9.3 Entering experimental data

This is done either by entering single equlibria with conditions and in addition using the command "enter experiment", see section 10.7 where the experimental data is given with an uncertainty. Each equilibrium with an experiment is given a unique name.

Often there are tables with values and instead of entering each of them there is a command "enter many\_equilibria", see section 10.9 with a simplified syntax.

When all equilibria with experiental data has been entered you have to give the command "set range", see section 26.16 to give the first and last equilibrium number that should be used in the assessment. If necessary this range can be extended during the assessment.

All the experimental data should also be entered as a mcro file to keep a documentation.

### 2.9.4 Saving the state of the assessment

Any time during an assessment it is possible to save the values of all assessed parameters and the calculated experimental equilibria by the command "save unform *filename*", see section 24.6. With this command the data inside OC will be written as an unformatted Fortran file and this can be saved and later read back into the OC software by the command "read unfomatted *filename*", see section 23.5. If these commands are inside a macro file prefix the filename with "./" to read and write on the same directory as the macro file.

These unformatted files are very convenient but beware that they may not be portable to other operating systems or even other versions of OC compiled with different Fortran compilers. It may change in future releases of the OC software. Thus keep printouts and macro files also if you later want to make modifications.

### 2.9.5 Performing the assessment

There are many decisions to make during the assessment and a general description how to perform an assessment can be found in the book by Lukas et al [07Luk]. It is never possible to try to assess all parameters using all experiments in a single step. Normally the user selects different sets of experimental data by the "set weight" command, see section 26.24 and fits a few model parameters to these using the command "set variable-coeff", see section 26.22. This can typically an enthalpy of mixing or a heat capacity function for a compound.

The command to run the least square fit is "optimize" followed by the maximum number of iterations, see section 20. If zero is given a single loop is made through all equilibria with nonzero weights within the specified range is made. It is also possible to use the command "calculate all", see section 7.1, to calculate all non-zero weight equilibria. With the latter command you can turn on the grid minimizer, in the optimize command the grid minimizer is always turned off.

When the optimize command is given with nonzero maximum there will be output on the screen at regular intervals giving the current values of the optimizing coefficients and the value

of the sum of squares. When the oprimization is finished there will also be a listing of the errors for all experiments.

With the command "list opt short", see section 16.10.8, the current values of the optimizing coefficients and all equiliria with the experimental data is listed together with the sum of squares. New selection of equilibria or weights can be made and the values obtained for the optimizing coefficients must also be reasonable but to know what is reasonable is not always easy. These steps are repeated until the user is satisfied or exhausted.

Macro files to calculate and plot of the calculated properties overlayed with the experimental data should be preoared and run regularly as just looking at numbers is not sufficient.

At a later stage solubilities and phase diagram data are used but in many cases reasonable guesses of the start values of model parameters must be made to be able to calculate the equilibrium with the experiment. Great care must be taken that the calculated equilibria for the inital model parameters are reasonably close to the experimental. Parts of the experimental phase diagram may have to be assessed separately and the metastable extrapolations of the different phases checked.

Sometimes a phase appears in a region where it should not be stable and additional fictitious experimental data may have to be added to prevent this to happen.

At the end the assessment should be written up and published.

# 2.10 Environment and startup macro file

The OC program will look for an environment variable called OCHOME and if it finds this it will look for a file start.OCM which will be executed before the user gets control. This can typically be useful to set some variables like the plot terminals, see section 10.8. If there is no OCHOME environment variable the "working directory" will be searched.

The ochelp.hlp file should be copied to this OCHOME directory.

### 2.11 User interface feedback

OC has grown organically and although the basic concepts has been quite clear the implementations of several of these has become rather confusing. This will eventually require some cleaning up of the user interface.

If you like a Graphical User Interface you are welcome to write it.

A central part of any thermodynamic software is the modeling of the phases. A new PDB format for databases may help a little with the specification of the models. An attempt has been made in this version to clean up the way a model is specified and used. At present you must first ENTER the phase to give a name, basic model, sublattices and constituents. Then use the AMEND command to add magnetism, a disordered fraction set and/or use BCC/FCC permutations. Originally some of these things were set by the command SET PHASE ... BIT

and that was not very clear.

Some computational options like for the grid minimizer are still set with several different commands. It is useful for the developers to have some feedback from users to organize this better.

# 3 All commands

The commands in alphabetical order as listed with the?

ABOUT	EXIT	MAP	SELECT
AMEND	FIN	NEW	SET
BACK	HELP	OPTIMIZE	SHOW
CALCULATE	HPCALC	PLOT	STEP
DEBUG	INFORMATION	QUIT	
DELETE	LIST	READ	
ENTER	MACRO	SAVE	

Many of the commands have "subcommands" and usually there is a default (listed within slashes //) which is selected by pressing return. You can type commands and subcommands and other parameters on the same line if one knows the order, using a comma, "," to select the default.

# 3.1 Options

There some options that can be set for the whole session or for just a single command. The options are identified by a / in front like /output=myfile.dat.

The options can be specifies directly after a command. Only a few are implemented.

- /OUTPUT=file name open a file and write. Note that if you have popup windows enabled this will open unless you type the file name on the same line as the command. In a macro file must prefix the file name with "./" to have the output (or append) on the same directory as the macro file. See also section 2.1.1 and 2.8.
- /APPEND=file name append output to a file, any previous content is kept.
- /ALL apply for all
- /FORCE override normal restrictions
- /VERBOSE write information while executing
- /SILENT do not write anything except fatal error messages

# 4 About

Some information about the OC software.

## 5 Amend

Intended to allow changes of already entered data. Only some of the subcommands are implemented.

BIBLIOGRAPHY EQUILBRIUM PARAMETER SYMBOL

COMPONENTS GENERAL PHASE TPFUN\_SYMBOL

CONSTITUTION LINE QUIT ELEMENT OPTIMIZING\_COEFS SPECIES

The default selection is PHASE.

# 5.1 amend Bibliography

The text for bibliographic reference identifier can be amended. The reference identifier is case insensitive.

# 5.2 amend Components

By default the elements are the components. This command can set any orthogonal set of species as components. The number of components cannot be changed by this command. The new components must exist as species and be orthogonal.

You must use components when setting conditions on amounts, mole fractions, chemical potentials or activities. Note that when you have other components than the elements you may have negative mole fractions and phase amounts.

### 5.3 amend Constitution

The program will ask for a phase name and you can set the amount and constitution of the phase. This will be used as initial constitution for a calculation unless the grid minimizer is used.

### 5.4 amend Element

The data for the element can be amended, not implemented yet.

# 5.5 amend Equilirium

Not sure what could be amended and anyway not implemented.

### 5.6 amend General

A number of user specific settings for defaults can be made:

- The name of the system.
- The level of the user (beginner, frequent user, expert). This may affect the behavior of the program (not implemented yet).
- If global minimization is allowed or not.
- If the grid minimizer is allowed to merge gridpoints in the same phase after global minimization.
- If the grid minimizer can automatic create composition sets is allowed or not.
- If redundant composition sets can be deleted automatically after an equilibrium calculaion.

Note that these and some other general feautures can also be changed by the command **SET BIT GLOBAL** 

### 5.7 amend Line

After a STEP or MAP command it is possible to give the command LIST LINE to list all calculated equilibria or AMEND LINE which allows you to EXCLUDE lines or INCLUDE lines from the plotting.

# 5.8 amend All optimizing coefficients

The values of the optimizing coefficients, see section 26.16 can be rescaled (start values set to current values) or recovered (current values set to previous start values).

### 5.9 amend Parameter

The possible parameters that can be amended depend on the model of the phase. By specifying a parameter you can change its expression. This is not yet implemented you must use the command **ENTER PARAMETER** to change the parameter expression.

# $5.10 \quad amend \; \mathrm{Phase} \; < phase \; name >$

You must first specify the phase name and then you can amend some of the properties of the phase:

ADDITION DEFAULT\_CONSTIT FCC\_PERMUTATIONS QUIT

AQUEUS\_MODEL DIFFUSION FLORY\_HUGG\_MODEL UNIQUAC\_MODEL

BCC\_PERMUTATIONS DISORDERED\_FRACS GADDITION

COMPOSITION\_SET FCC\_CVM\_TETRAHDR QUASICHEM\_MODEL

### 5.10.1 amend phase < ... > Addition

Additions are used to give a contribution to the Gibbs energy of a phase using more or less physically based model. Usually they require additional model parameters, see section refsc:paramid. The difference between addition and other things that can be amended may not always be very clear. The possible additions are

CRYSTAL\_BREAKDWN LOW\_CP\_MODEL QUIT SMOOTH\_CP\_STEP ELASTIC\_MODEL\_1 MAGNETIC\_CONTRIB SCHOTTKY\_ANOMATY TWOSTATE\_LIQUID

BEWHERE! The OC software allows you to mix many types of additions for a phase but it is up to the user to defend the physical reasons for this!

### 5.10.1.1 amend phase ... addition Crystal\_breakdwn

When extrapolating the Gibbs energy of the solid phases above the melting T the heat capacity of the metstable solid is simply extrapolated from the solid range this can lead to extremely high heat capacity values which may make the solid stable again. It is also a problem when using the Kopp-Neuman rule to model the heat capacity of solution phases as well as compounds which are stable above the melting T of the element.

In the 1991 SGTE unary database a breakpoint was introduced at the melting T of the element but that gives the false impression something happends in the solid phase at the melting T.

Introducing a crystalline breakdown T (CBT), which should be several 100 K above the melting T, is a way to handle this but the actual function is still discussed.

### 5.10.1.2 amend phase ... addition Elastic\_model\_1

A contribution to the Gibbs energy due to elastic strain can be added. This also requires values of the elastic constants and lattice parameters, see section 2.6.1.

There is no code to calculate the elastic energy implemented yet.

### 5.10.1.3 amend phase ... addition LowT\_Cp\_model

The Einstein model for heat capacities from 0 K has been implemented. It requires a value of the property Einstein T as listed in section 2.6.1.

### 5.10.1.4 amend phase ... addition Magnetic\_contrib

The Inden-Hillert and the modified Inden-Qing model for the magnetic contribution to the Gibbs energy can be set by this command This depends on model parameters describing the Curie and Neel temperatures and the Bohr magneton number, as listed in model parameters identifiers 2.6.1, for the phase.

The Qing-Xiong model is selected by giving zero (0) for the question about the anti-ferromagnetic factor.

The Inden-Hillert model is described in [07Luk] and the Inden-Qing-Xiong modified model requires separate values of the Curie and Neel Temperatures and either an "effective" Bohr magneton number or individual Bohr magneton numbers for the constituents of the phase.

### 5.10.1.5 amend phase ... addition Quit

You did not really wanted to add any addition.

### 5.10.1.6 amend phase ... addition Schottky\_anomalty

Some physical phenomena can increase the heat capacity for a phase and this addition can describe this. It uses two model parameter identifiers, TSCH and CSCH that may depend on the composition. TSCH specify the T for the anomality and CSCH the maximum contribution to the heat capacity (J/mol/formula unit).

### 5.10.1.7 amend phase ... addition Smooth\_Cp\_step

The 3rd generation thermodynamic databases extrapolate to 0"K and require that the heat capacity is zero at 0 K. This means it is impossible to use  $T * \ln(T)$  terms (and also negative powers of  $T^{-n}$ ) but there may be some physical phenomena that causes an incremental increase of the heat capacity at some temperature. Ignoring the physical reason for such an increase this "smooth\_ $C_P$ \_step addition will provide such this using two parameters, THT2 to specify T and DCP2 to specify the increment in heat capacity. It uses the same mathematical expression as the Einstein heat capacity function.

### 5.10.1.8 amend phase ... addition Twostate\_liquid

The two-state model for the hear capacity for the undercooled liquids can be added. It assumes a low T amorphous state modeled as an Einstein solid and requires an Einstein T. For the liquid transition it uses the model\_parameter\_ident bf G2, both of which are listed in section 2.6.1.

You must specify parameters for THET and G2 for all constituents of the phase and possibly also interaction parameters to specify the composition dependence.

The implementation of this addition is not finished.

### 5.10.2 amend phase ... Aqueous\_model

A model with dilute configurational entropy. Not implemented yet.

### 5.10.3 amend phase ... BCC\_permutations

This is intended for the 4 sublattice CEF model for BCC ordering. Due to crystallographic symmetry several model parameters must be identical such as

$$G(BCC,AL:FE:FE:FE)=G(BCC,FE:AL:FE:FE)=G(BCC,FE:FE:AL:FE)=G(BCC,FE:FE:AL)$$

and this command means these parameters need to be entered only once. This affects the data storage and the calculation of the Gibbs energy is slightly more efficient. The same applies for the FCC\_permutations but the BCC tetrahedron is slightly asymmetric which makes it a bit more complicated than the FCC. There can be a 5th sublattice with interstitials.

### 5.10.4 amend phase ... Composition set

Each phase has by default a single composition set. If the same phase can exist as stable (or metastable) with two or more compositions (miscibility gaps or order/disorder transformations) you may have to amend the phase by creating additional composition sets.

Composition sets can also be created automatically by the grid minimizer during an equilibrium calculation. In such a case the composition set will have the suffix \_AUTO,

Composition sets of a phase can be created and deleted. Phases with miscibility gaps or which can exist with different chemical ordering like A2 and B2 must be treated as different composition sets. You can specify a prefix and suffix for the composition set. Extra composition sets will always have a suffix #digit where digit is a number between 2 and 9. You cannot have more than 9 composition sets.

The composition set number is given after the phase name and preceded by a hash character #. In the OCASI interface and some more cases phase tuples are used to identify a phase and a composition set by a single number. As composition sets can be created and deleted a phase

tuple index for the 2nd or higher composition set may change between calculations.

In some cases it may be interesting to calculate metastable states inside miscibility gaps and you can prevent the automatic creation of composition sets by turning off the global minimazation using AMEND GENERAL or for an individual phase by SET PHASE ... BIT NO\_AUTO\_COMP\_SET

### 5.10.5 amend phase ... Default Constitution

The default constitution of a phase can be set. Unless the grid minimizer is used this will be used for the first calculation with the phase and sometimes if there are convergence problems. NOTE that if you want to specify a default constitution for the second or higher composition set of a phase you must specify the composition set with the phase name!

Depending on the minimizing software used the initial constitution can be important to find the correct equilibrium if the phase has ordering or a miscibility gap.

For each constituent you can specify a minimum > or maximum < fraction or give NONE if there are no default.

If a phase has miscibility gaps and you have created composition sets with default constitutions the grid minimizer will try to select the composition set with a composition closest to the default for a stable phase.

To temporarily set a new constitution of a phase use the command AMEND CONSTITUTION <phase> or CALCULATE PHASE ... .

### 5.10.6 amend phase ... Diffusion

This is to specify how the diffusion coefficient matrix should be calculated when simulating a phase transformation. Normally the mobilities for the constituents of the phase are read from the database but you may use different "depended" and "independent" constituents in the diffusion model and also some other factors. This command is intended for such use. It is not implemented yet.

There is no intention that OC itself should simulate diffusion but as the diffusion coefficients are strongly dependent on the thermodynamic factor (the Darken stability matrix) which represent the second derivatives of the Gibbs energy it is convenient to include some properties used in a simulation in the thermodynamic software.

### 5.10.7 amend phase ... Disordered fraction sets

For phases with several sublattices the Gibbs energy of the phase can be divided into two sets of fractions where the second or "disordered" set have only one or two sublattices and the fractions on these represent the sum of fraction on some or all of the first or "ordered" set of sublattices.

This is particularly important to model the Gibbs energy for phases with ordering like FCC, BCC and HCP and for intermediate phases like SIGMA, MU etc.

### 5.10.8 amend phase ... FCC\_CVM\_tetradrn

This model is intended for the CVM tetrahedron model for FCC and HCP. Not implemented yet.

### 5.10.9 amend phase ... FCC\_permutations

This is intended for the 4 sublattice CEF model for FCC ordering. Due to crystallographic symmetry several model parameters must be identical such as

$$G(FCC,AL:FE:FE:FE)=G(FCC,FE:AL:FE:FE)=G(FCC,FE:FE:AL:FE)=G(FCC,FE:FE:FE:AL)$$

Setting this means that unique model parameters need to be entered only once, the software will take care of all permutations. HCP permutations are also handled with this command as the HCP tetrahedron model is identical to the FCC. There can be a 5th interstitial sublattice.

### 5.10.10 amend phase ... Flory\_Huggins model

A model when the configurational entropy takes into account that the constituents have very different size. Not implemented yet.

### 5.10.11 amend phase ... Gaddition

You can add a constant value of the Gibbs energy to a phase in Joule per formula unit. This is a simple way to implement a for example a nucleation barrier.

### 5.10.12 amend phase ... Quasichemical

There are several quasichemical models for the liquid that only describes the short range ordering (SRO). None of them are yet implemented.

### 5.10.13 amend phase ... Quit

Do not amend anything for the phase.

### 5.10.14 amend phase ... UNIQUAC

The UNIQUAC model for polymers is on its way to be implemented.

# 5.11 amend Quit

Do not amend anything (more).

# 5.12 amend Species

Not implemented yet.

# 5.13 amend Symbol

This command is a bit special. It is mainly used in assessments to specify that a particular symbol must not be evaluated except when specified explicity, or when calculating a specific equilibrium.

The main reason is that a symbol can refer to another symbol and thus all symbols are normally evaluated whenever the value of any symbol is requested. This is to ensure that all symbol values are consistent and refer to the same calculated equilibrium. But in certain cases you may want to enter a symbol that is only evaluted when referenced explicity or at a specific eqilibrium and this can be set with this command.

When you want to compare the value of a thermodynamic property, like the enthalpy, in two equilibria you must be able to store the calculated enthalpy from one equilibrium in a symbol. For example if you have experimental data on the heat difference for a compound at various T. In such a case the enthalpy at the reference T can be stored in a symbol, which has been amended with this command to specify at which equilibrium it should be evaluated. In all other equilibria the value of this symbol will have the value at the specified equilibrium. See also the documentation on the assessment procedure, section 2.9.5.

You cannot amend the expression for the symbol except symbols that represent a constant value.

# 5.14 amend Tpfun\_Symbol

You can replace a TP function with a new expression.

# 6 Back

Return back from the command monitor to the application program. In the OC software itself it means terminate the program.

# 7 Calculate

Many different things can be calculated. The normal thing to calculate is **equilibrium**, the other things are special.

ALL\_EQUILIBRIA NO\_GLOBAL SYMBOL WITH\_CHECK\_AFTER

EQUILIBRIUM PHASE TPFUN\_SYMBOLS GLOBAL\_GRIDMIN QUIT TRANSITION

# 7.1 calculate All equilibria

Intended for the assessment procedure. Calculates all equilibria with non-zero weight as set by the command **SET RANGE**. It can also be used for other purposes, for example testing the parallelization. The equilibria can be entered by the command **ENTER MANY\_EQUILIB**.

# 7.2 calculate Equilibrium

The normal command to calculate the equilibrium of a system for the current set of conditions and phase status. You can calculate a metastable equilibrium if some phases that should be stable have been set dormant or suspended or if automatic creation of composition sets is not allowed. If conditions allow the grid minimizer is used to find start values unless the grid minimizer is explicitly turned of.

Before this command you must have entered thermodynamic data from a database or interactively and used the command **set condition** to set as many conditions as you have components plus two. The command **set status phase** and **set input amount** can also be used to set conditions.

### 7.3 calculate Global Gridmin

Calculate with the global grid minimizer without using this result as a start point for the general minimizer. Used to debug the grid minimizer.

### 7.4 calculate No\_Global

Calculate the equilibrium without using a global grid minimizer to generate start constitutions. The current equilibrium is used as start point. Can be quicker when only small changes of conditions made since previous calculation and this is how equilibria is calculated during STEP and MAP. It means no check of new miscibility gaps.

# 7.5 $calculate \ Phase < phase \ name >$

You must provide a phase name, the amount of the phase and if you should use the current constitution or enter a new.

The Gibbs energy of a phase and possible derivatives and some other things can be calculated. Mainly for debugging the implementation of models and testing the software.

ALL\_DERIVATIVES DIFFUSION\_COEFF ONLY\_G CONSTITUTION\_ADJ G\_AND\_DGDY

### 7.5.1 calculate phase ... All\_Derivatives

The Gibbs energy, all T and P derivatives and all first and second derivatives with respect to constituents for the specified phase for current T, P are calculated and listed.

### 7.5.2 calculate phase ... Constitution\_Adjust

You will be asked to enter a new composition (note NOT the constitution and default is current) of the phase and this command will then calculate the Gibbs energy and all chemical potentials. For a phase with sublattices the constitution of the phase will be adjusted to have the minimum Gibbs energy for the given composition.

It is useful when one or more components are parts of several constituents, for example in a gas or for phases with order/disorder transitions, in particular when the corresponding subroutine is used in simulations.

### 7.5.3 calculate phase ... Diffusion\_Coefficients

You will be asked to enter a new composition (default is current) of the phase and this command will then calculate the Darken stability matrix

$$\frac{\partial^2 G}{\partial N_{\rm A} \partial N_{\rm B}}$$

for all components (see the documentation of the minimiser) and also all mobility values (if there are any).

### 7.5.4 calculate phase ... G\_and\_dGdy

The Gibbs energy, all T and P derivatives and all first derivatives with respect to constituents for the specified phase for current T, P are calculated and listed.

IMPORTANT NOTE: The value of  $\frac{\partial G_m}{\partial y_i}$  is NOT the chemical potential,  $\mu_i$  of component i. The understanding of thermodynamics is often poor and the user is reminded that the chemical

potential of a component i is defined as:

$$\mu_i = \left(\frac{\partial G}{\partial N_i}\right)_{T,P,N_{i \neq i}}$$

where G is the integral Gibbs energy and all  $N_i$  are independent variables. When we model the molar Gibbs energy,  $G_m$  as a function of the constituent fractions,  $y_i$ , these fractions are not independent and for a substitutional model, where  $y_i = x_i$  i.e. the mole fractions, the chemical potential is calculated from  $G_m$  using:

$$\mu_i = G_m + \left(\frac{\partial G_m}{\partial x_i}\right)_{T,P,x_{j\neq i}} - \sum_j x_j \left(\frac{\partial G_m}{\partial x_j}\right)_{T,P,N_{k\neq j}}$$

because the mole fractions,  $x_i$  are not independent.

#### 7.5.5 calculate phase ... Only\_G

The Gibbs energy and all T and P derivatives calculated and listed for the specified phase for the current values of T, P.

### 7.5.6 calculate phase ... Quit

Do not calculate anything for the phase.

## 7.6 *calculate* Quit

Do not calculate anything at all.

# 7.7 calculate Symbol

A state variable symbol or function is calculated using the results from the last equilibrium or grid minimizer calculation. It is used in particular for calculation of "dot derivatives" like H.T for the heat capacity.

If a wildcard, "\*", is given as name all symbols, except dot derivatives and symbols that must be specified explicity and those that should be calculated for another specified equilibria. See section 5.13.

# 7.8 calculate Tpfun\_Symbols

All or a specific TPFUN symbol are calculated for current values of T and P.

### 7.9 calculate Transition

After calculating an equilibrium you can calculate directly when a phase will appear or disappear by releasing one of the conditions you have specified. Typically this is used to calculate the melting temperature of an alloy or a solubility limit.

You specify the phase name and the condition to be released. The program will set this phase as FIXED with zero amount and remove the condition you specified and calculate the equilibrium. The calculation may fail if the phase cannot be set stable with zero amount. If successful the removed condition will be set to the value calculated and the phase set stable with zero amount.

# 8 Debug

Several possibilities to trace calculations will be implemented in order to find errors but very little is working yet.

## 8.1 debug Elasticity

Intended to test the model for strain and stress. Not implemented.

# 8.2 debug Free lists

Only for experts.

# 8.3 debug Stop\_on\_Error

The program will stop at the command level after printing the error message if an error has occurred when using macro file. This should make it easier to to find errors occurring when running macro files.

However, it is not implemented.

# 9 Delete

It is quite difficult to delete anything when the data structure is so involved. In many cases it may be better to enter the data again without the data that should be deleted. But there are a few things that must occationally be deleted.

COMPOSITION\_SET EQUILIBRIUM QUIT STEP\_MAP\_RESULTS ELEMENTS PHASE SPECIES

## 9.1 delete Composition set

The first composition set of a phase cannot be deleted. Otherwise there is usually no problem unless several equilibria are entered because the composition set must be deleted in all equilibria. Composition sets are created and deleted during normal equilibrium calculations to detect miscibility gaps.

### 9.2 delete Element

Dangerous and will probably never be implemented.

## 9.3 delete Equilibrium

Dangerous but sometimes necessary. Done automatically at a second STEP or MAP command if you specifies to delete previous results.

### 9.4 delete Phase

Dangerous and will probably never be implemented.

## 9.5 delete Quit

Do not delete anything.

# 9.6 delete Species

Not implemented yet and will probably never be.

# 9.7 $delete Step\_Map\_Results$

This removes all equilibria and saved equilibria associated with STEP and MAP commands. It also deletes the axis.

# 10 Enter

In most cases data will be read from a database file. But it is possible to enter all thermodynamic data interactively. This should normally start by entering all elements, then all species (the elements will automatically also be species) and then the phases.

A species have a fixed stoichiometry and possibly a charge. The species are the constituents of the phases.

A phase can have sublattices and constituents and also various additions like magnetic, low T heat capacity etc. which are specified by the **AMEND** command efter entering the phase (but normally before any model parameters for the phase are entered).

TPFUN symbols can be used to describe common parts of model parameters. See section 2.6.3 for an explation.

Each model parameter of a phase is entered separately. You may use TPFUN symbols which are already entered.

At present the multicomponent CEF model and the ionic 2-sublattice liquid model are the only basic models implemented. The CEF model includes as special cases the gas phase, regular solutions with Redlich-Kister Muggianu model and phases with up to 9 sublattices and ionic constituents. These models describe the basic configurational entropy contribution to the phase, models such as the magnetic contribution and low T heat capacity can be added to a phase with the **AMEND** command.

The enter command is also used to enter bibliographic data, equilibria for assessments and many other things.

The subcommands are:

BIBLIOGRAPHY	EQUILIBRIUM	OPTIMIZE_COEFF	SPECIES
COMMENT	EXPERIMENT	PARAMETER	SYMBOL
CONSTITUTION	GNUPLOT_TERMINAL	PHASE	TPFUN_SYMBOL
COPY_OF_EQUILIB	MANY_EQUILIBRIA	PLOT_DATA	
ELEMENT	MATERIAL	QUIT	

# 10.1 enter Bibliography

Each model parameter must have a bibliographic reference to ensure everyone can find the origin of its value. When entering a parameter a bibliographic reference symbol is given and with this command you can give a full reference text for that symbol like a published paper, a report or simply the reason for the value. The date and name of the responsible should also be given.

#### 10.2 enter Comment

A line of comment text can be added to an equilibrium. It is particularly important when entering experimental data to give the reference to the data.

### 10.3 enter Constitution

The constitution (fraction of all constituents) of a phase can be entered. This is a way to provide start values for an equilibrium calculation (when not using grid minimizer). To calculate the Gibbs energy for a specific phase at a specific constitution use the command **CALCULATE PHASE**.

## 10.4 enter Copy of equilibrium

This command creates a copy of the current equilibrium with the same set of conditions and related data.

#### 10.5 enter Element

The data for an element is entered. It consists of is symbol, name, reference state, mass, H298-H0 and S298. The latter two values are never used for any calculation.

## 10.6 enter Equilibrium

You can have several equilibria each with a unique set of conditions including phase status (dormant, suspended, fix or entered) but all with the same components and thermodynamic data. This is useful for compare different states, to simulate transformations and to assess model parameters as each experimental or theoretical information represented as an equilibrium.

All equilibria use the same thermodynamic data but they have an independent set of conditions and result data structure and they can be calculated in parallel.

# 10.7 enter Experiment

This is used for assessments, experimental data can be specified for an equilibrium. The experiment is a state variable or symbol which can be set equal to the experimental value followed by a colon, ":" and its uncertainty.

In some cases an experimental value can be an upper or lower limit. In such cases the ">" or "<" can be used. The value of the uncertainty will then be interpreted as a penalty factor if the calculated value is outside the specified limit.

### 10.8 enter GNUPLOT Terminal

For plotting OC generates a command file for the GNUPLOT [GNUPLOT] software. GNU-PLOT can be downloaded free for most OS but depending on your screen and other hardware you may prefer to specify your prefered set of terminals. On Windows the defaults are:

1 SCREEN > set terminal wxt size 900.600

2 PS > set terminal postscript color solid

3 PDF > set terminal pdf color solid size 6,4 enhanced fontscale 0.45

4 GIF > set terminal gif

The text after the > is written on the GNU command file. You can change these or add additional terminals. You can also change these in the source code (userif/pmon6.F90 file) or use a macro file OCHOME/start.OCM file to set them.

## 10.9 enter Many Equilibria

This command is intended for adding tables of experimental data of the same type. It can also be used for calculation of many equilibria using the **calculate all** command. The user first enters a TABLE HEAD giving the necessary phase status, conditions, experiments etc. In this "head" some values of text can be referred to columns in the following table using the "@" character followed by a digit 1 to 9, where the digit is the column number.

The prompt for input to the table head is "table head::"

In the examples below, taken from the parallel 2.OCM macro file, user input is **in bold** and explanations in italics.

• By default all phases are suspended so the user must forst specify the phases with dormant, entered of fixed status (including amount) like

Table head:: entered 0 \* all phases should be entered

Table head:: fix 0 liquid liquid should be fix with 0 moles

Table head:: fix 1 @2 the phase in column 2 should be fix with 1 moles

• The conditions can be given using the @ character to indicate vaules that are given in the specified column in table to follow.

Table head:: condition t=@1 p=1e5 n=1 w(cr)=@3 w(mo)=@4

• Optional calculations of entered symbols

Table head: calculate cp

• Optional listing of state variables

Table head: list HM tc(bcc)

• Optional experimental data

Table head: experiment x(liquid,cr) = @5:.01, x(bcc,cr) = @6:.02

• Optional reference state

The reference state for a component can be set.

Table head: reference O gas \* 1e5

The reference state for the component O will be gas at the current T and 1 bar.

• Optional plot\_data specifying a dataset number and coordinates to be plotted and a symbol. The coordinates can be table columns. Use the dataset numbers to have data of the same type together like enthalpies, phase diagram data etc.

Table head: **plot 1 @1 @2 5** 

• Optional comment

Table head: comment experimental data from Kubaschewski 1955

• The table head is finished by an empty line or "table\_start"

For the rows in the table the user must first provide a unique name for each equilibrium (that is counted as column 0 (zero)) and values for all columns referenced in the table head like:

Table row: EQ1 1573 BCC 0.3 0.05 0.12 0.28 Table row: EQ2 1623 BCC 0.3 0.10 0.18 0.24

The table is finished by an empty line or

Table row: table\_end

### 10.10 enter Material

The user will be asked for a name of the material and possibly a database. Then he can give elements and their amount in mass percent or mole fraction. Finish with an empty line. Then he must specify the temperature and the program will automatically make a calculation at 1 bar. For example:

```
OC4: enter mat
Database:steel7
Elements: C , MO, V , CR, FE, SI,
Major element or material:fe
Input in mass percent? /Y/:
Input expected in mass percent
First alloying element:c
Mass percent: /1/:
Second alloying element:cr
Mass percent: /1/: 5
Third alloying element:mo
Mass percent: /1/: 8
Next alloying element:v
Mass percent: /1/:
Next alloying element:
 3E reading a TDB file
 3D em: W''_{N}(C)=1 W''_{N}(CR)=5 W''_{N}(MO)=8 W''_{N}(V)=1
                                                   N=1
```

```
Temperature /1000/:
```

3Y Constitution of metastable phases set

3Y Composition set(s) created:

Gridmin: 18846 points 6.25E-02 s and 78 clockcycles, T= 1000.00

Phase change: its/add/remove: 5 0 21

Equilibrium calculation 19 its, 6.2500E-02 s and 94 clockcycles

The user can specify another composition of the same alloy with the same command or use other commands such as **SET CONDITION** and **CALCULATE** or calculate diagrams using **SET AXIS**.

## 10.11 enter Optimizing coefficient

The number of TP symbols for the coefficients to be optimized are entered. They have the names A00 to A99. They are used in model parameters and can be varied by the optimization procedure to minimize the difference between the experimental data and the same property calculated from the models of the phases. You can also specify the size of the workspace needed for the optimization. The default value, 2500, is usually sufficient.

### 10.12 enter Parameter

A model parameter is defined by its identifier, the phase and constituent array and the degree. A parameter can be a constant or depend on T and P. The parameter will be multiplied with the fractions of the constituents given by its constituent array. See the documentation of the GTP model package or the book by Lukas et al[07Luk] for more information about thermodynamic models.

For example G(LIQUID,CR) is the Gibbs energy of liquid Cr relative to its reference state, normally the stable state of Cr at 298.15 K and 1 bar, and called an endmember.

For a gas molecule the parameter G(GAS,C1O2) is also an endmember and represent the Gibbs energy of the C1O2 molecule relative to the reference states of C (carbon) and O (oxygen).

For interaction parameters the components are separated by a comma "," as in G(LIQUID, CR, FE).

For phases with sublattices the constituents in each sublattice are separated by a colon, ":" and interacting constituents in the same sublattice by a comma, ",". For example: G(FCC,FE:C,VA) is the interaction between C (carbon) and VA (vacant interstitial sites) in the FCC phase.

The only binary excess model implemented in OC is the Redlish-Kister with the Muggianu ternary extrapolation method.

$$L_{A,B} = \sum_{\nu=0}^{n} {}^{\nu}L_{A,B}(y_{A} - y_{B})^{\nu}$$

where the degree,  $\nu$ , of the interaction parameter is specified after a semicolon, L(phase,A,B; $\nu$ ).

For ternary parameters and for reciprocal parameters the Hillert model for composition dependence is implemented, see [07Luk].

You can store many different types of data in OC with different parameter identifier. Some of the parameters are not related to the thermodynamic properties but as they depend on the phase, T, P and composition it is convenient to store them together with the thermodynamic data. For example the mobility of Fe in BCC (including an empty interstitial sublattice) is specified as: MQ&FE(BCC,FE:VA).

An explanation of the identifiers implemented in OC can be found in section 2.6.1. The current list can be obtained by the command **LIST MODEL\_PARAM\_ID**. All of them can be composition dependent. Some cannot depend on T or P or neither. Many kinds of the parameters are available but in some cases the software for the models to handle them are not implemented. The value of a model parameter can be obtained using **LIST MODEL\_PARAM\_VAL** or simply **SHOW**. You must specify phase and endmember for the parameter.

### 10.13 enter Phase

All thermodynamic data are connected to a phase as defined by its parameters, see **enter parameter**. A phase has a name with can contain letters, digits and the underscore character. It must start with a letter.

After the phase name you must specify a model. The model specification is implemented in a rather rudimentary way. The only recognized models are

- IDEAL for a single lattice phase without interactions (like GAS)
- RKM for a substitutional phase with interactions (like metallic liquid)
- I2SL for the ionic liquid phase (2 sublattices with variable site ratios). If the phase name is IONIC\_LIQUID this prompted as the default model.
- CQC means the "Corrected Quasichemical model" for liquids.
- CEF for any other phase with two or more sublattices

This list may be extended in a future version of OC. Many other model features like magnetism, quasichemical etc are specified with the **AMEND PHASE** command, see section 5.10. The AMEND PHASE command is also used to specify disordered fraction set, low temperature CP model and many other things.

For most models OC will ask for the number of sublattices and a phase can have 1 to 9 sublattices and you must specify the number of sites on each. Preferably use small integer values, if fractions are used at least 6 digits should be provided.

For some models, like the ionic liquid model, the number of sites may change with the composition of the phase so the number specified is irrelevant. See the book by Lukas et al [07Luk] for more details on models.

For each sublattice you must specify the constituents on the sublattice. A constituent that is not an element must already have been entered as a species, see section 10.16.

The **AMEND PHASE** command 5.10 is used for some additional model features like magnetism or permutations.

### 10.14 enter Plot data

This is used for assessments when combining experimental data in single equilibria with those in tables entered with the command "MANY\_EQUILIBRIA".

You can add points to be plotted from a single equilibrum to a dataset 1 to 9. The dataset must already exist as a file already opened by a command ENTER MANY\_EQUILIB.

### 10.15 enter Quit

Quit entering things.

# 10.16 enter Species

A species consists of a name and a stoichiometric formula. It can have a valence or charge. The name is often the stoichiometric formula but it does not have to be that. Examples:

- enter species water h2o
- enter species c2h2cl2\_trans c2h2cl2
- enter species c2h2cl2\_cis c2h2cl2
- enter species h+ h1/--1

There can be a problem with ambiguity with a species name like h20 if there is also a species h202. In such cases use a final unity, i.e. h201.

Single letter element names must be followed by a stoichiometric factor unless it is the last element when 1 is assumed. Two-letter element names have by default the stoichiometric factor 1.

- enter species carbonmonoxide c1o1
- enter species cobaltoxide coo

• enter species carbondioxide c1o2

The species name is important as it is the name, not the stoichiometry, that is used when referring to the species elsewhere like as a phase constituent. It is of course convenient to choose a species name similar to its stoichiometric formula but as shown above, that is not always sufficient.

Species symbol:

The symbol must start with a letter a A-Z and contain just letter, digits and the special characters "\_" (underscore), "\_" (minus) and "+" (plus).

Species stoichiometry:

The stoichiometry must contain element symbols followed by a stoichiometry factor. The stoichiometry factor 1 can be omitted for two-letter element symbols. The charge is given as "/-" or "/+" followed by a stoichiometry factor.

## 10.17 enter Symbol

The OC package has both "symbols" and "tpfun\_symbols", the latter has a very special syntax and can be used when entering parameters.

The symbols are designed to handle relations between state variables, you can define expressions like

### enter symbol KLBCR = X(LIQUID,CR)/X(BCC,CR);

where KLBCR is set to the partition of the Cr mole fractions between liquid and bcc.

The symbols also include "dot derivatives" like H.T which is the second derivative of the Gibbs energy with respect to the for the current system at the given set of conditions.

#### enter symbol CP = H.T;

If T and P are conditions and all other conditions are mass balance conditions CP is the heat capacity. It also takes account of the change of configurational entropy.

Currently H.T is the only dot derivatives allowed but more will be added as soon as possible.

# 10.18 enter Tpfun\_Symbol

This symbol is an expression depending on T and P that can be used when entering parameters. A TPfun can refer to another TPfun.

The program requests a name, if the symbol should be a function, constant or a table (tables not implemented).

If it is a function you must specify a low T limit, and expression consisting of simple terms (signed coefficients multiplied with T and P possibly raised to powers). A term may also be

multiplied with another TP function or by LN(termm) for the natural logarithm or EXP(term) for the exponential function of "term".

The "term" inside the parenthesis of an LN or EXP you may refer to another TP function or it can be a coefficient multiplied with powres of T or P.

It is not allowed to enclose terms by parenthesis.

The expression is terminated by a semicolon followed by an upper T limit and possibly Y meaning another expression above this T limit. The last T-range must be followed by N and a bibliographic reference.

TPFUNs have a strict syntax because the software must be able to fast calculate first and second derivatives with respect to T and P during equilibrium calculations, see section 2.6.3.

## 11 Exit

Terminate the OC software in Swedish, Ha en bra dag.

## 12 Fin

Terminate the OC software in French, Au revoir.

# 13 Help and?

Can give a list if commands or subcommands or parts of this help text.

## 14 HPcalc

Start the reverse polish calculator.

# 15 Information

The intention is to provide the on-line user who does not like to read manuals with additional explanations but not implemented yet.

## 16 List

Many things can be listed. Output is normally on the screen unless it is redirected by the /output=file name or /append=file nameoption.

AXIS	ERROR_MESSAGE	PARAMETER	STATE_VARIABLES
BIBLIOGRAPHY	LINE_EQUILIBRIA	PHASE	SYMBOLS
CONDITIONS	MODEL_PARAM_ID	QUIT	TPFUN_SYMBOLS
DATA	MODEL_PARAM_VAL	RESULTS	
EQUILIBRIA	OPTIMIZATION	SHORT	

### 16.1 list Axis

Lists the axis set by you.

## 16.2 *list* Bibliography

List the bibliographic references for the data.

### 16.3 *list* Conditions

Lists the current set of conditions set by you. If the degrees of freedoms are zero you can calculate an equilibrium.

### 16.4 list Data

Lists all thermodynamic data. The default is on SCREEN but you can also choose among the formats: LaTeX, MACRO, PDB and TDB.

The only formats implemented at present are SCREEN, PDB and TDB.

#### 16.4.1 list data LaTeX

The thermodynamic data will be formatted according to LaTeX for later inclusion in publications. Not implemented.

#### 16.4.2 list data Macro

The thermodynamic data will be written as a macro file that can later be read back into the OC software. Not implemented.

#### 16.4.3 list data PDB

A "Portable phase related Data Format" similar to the TDB file format adapted for OC.

#### 16.4.4 list data TDB

A variant of the TDB file format with Thermo-Calc flavor.

## 16.5 *list* Equilibria

Lists the equilibria entered. To list the results of the calculation of an equilibrium use **list** result.

## 16.6 *list* Error message

The message associated with an error code generated by OC can be listed

## 16.7 list Line equilibria

Lists the equilibria calculated during STEP or MAP commands. See also the command **AMEND LINE-EQUILIBRIA**.

# 16.8 list Model parameter identifiers

Lists the model parameter identifiers available in the current version of OC, see section 2.6.1.

# 16.9 list Model parameter value

The current value of a model parameter identifier can be listed. Note that the value is always phase dependent and may also depend on the composition set.

# 16.10 list optimization

Lists results of an optimization, several sub-options will be implemented but currently there is a short version only.

COEFFICIENTS DEBUG GRAPHICS MACRO CORRELATION\_MTRX EXPERIMENTS LONG SHORT

#### 16.10.1 *list optimization* coefficients

This gives a list of the coefficients and their values.

#### 16.10.2 list optimization debug

Not implemented yet.

#### 16.10.3 *list optimization* correlation\_matrix

Not implemented yet.

### 16.10.4 list optimization experiments

List of experiments in the equilibria with non-zero weights.

### 16.10.5 list optimization graphics

A figure with the experimental values on the X axis and calculated values on the Y axis for all experiments. Not implemented yet.

#### 16.10.6 list optimization long

Not implemented yet

### 16.10.7 list optimization macro

A listing of all thermodynamic data and current values of model parameter and experimental data with current weight. This can be read back as a start of a re-assessment and an important documentation of the current state of the assessment. But not yet implemented.

#### 16.10.8 list optimization short

This specifies the data and hour of the listing and first a table with the optimizing coefficients with name, current value, start value, scaling factor and its relative standard deviation.

In the first table all the optimizing coefficients with non-zero values are listed together with the current values, the start values and their scaling factor (usually the same as the start value). In the column "RSD" the Relative Standard Deviation" should appear but it is not yet calculated correctly. Last column is the name of the TP symbol(s) where the coefficient is used.

After that all equilibria with non-sero weights are listed together with their experimental data, both the prescribed value, the uncertainy and the currently calculated one. In the last column the error is listed.

Listing of optimization results: date 2018.08.20 : 12h47

List of coefficients with non-zero values

Name	Current value	Start value	Scaling factor	RSD	Used in
A11	3.46818E+02	4.00095E+02	4.00095E+02	1.25070E-06	_GFCCABO
A12	-5.66234E+01	-6.52871E+01	-6.52871E+01	1.33802E-06	
A13	-2.10028E-02	-1.30393E-02	-1.30393E-02	8.97167E-06	_GFCCABO

List of 4 equilibria with 8 experimental data values

No Equil name	Weight Experiment \$ calculated	Error
2 FCC1_ZA	1.00 SM=17:1 \$ 17	9.8995E-09
2	1.00 CP1=18:1 \$ 17.28685	7.1315E-01
3 FCC2_ZB	1.00 HDIFF=9000:500 \$ 9997.813	-1.9956E+00
3	1.00 CP1=20:DCP \$ 22.55698	-2.5570E-02
4 FCC3_ZC	1.00 HDIFF=15000:500 \$ 14719.24	5.6152E-01
4	1.00 CP1=22:DCP \$ 24.65726	-2.6573E-02
5 FCC4_ZD	1.00 HDIFF=20000:500 \$ 19860.72	2.7856E-01
5	1.00 CP1=24:DCP \$ 26.75754	-2.7575E-02

Final sum of squared errors: 4.88614E+00 using 8 experiments and 3 coefficient(s). Degrees of freedom: 5, normalized error: 9.7723E-01

In the list of equilibria with non-zero weight the first column is a sequential equilibrium number assigned by the software. Then the name of the equilibrium assigned by the user. The third column is the weight, only equilibria with nonzero weight are listed. Then comes a column with the experimental property and value and after the dollar sign its calculated value with the present set of coefficients. The rightmost column gives the difference for each experiment  $i, q_i$  that should be as close to zero as possible:

$$q_i = \frac{z_i^{\text{exp}} - z^{\text{calc}}}{\sigma_i} w_i \tag{1}$$

where  $i, z_i^{\text{exp}}$  is the experimental property,  $z_i^{\text{calc}}$  is the same property calculated from the model and  $\sigma_i$  is the experimental uncertanty and  $w_i$  is the weight assigned to equilibria with the experiment. If  $w_i = 1$  and  $q_i$  is between -1 and 1 the experiment has been fitted within the experimental uncertanty.

The least square routine tries to determine coefficients to make the sum of all  $q_i^2$  as small as possible.

At the end of the listing  $\sum_i q_i^2$  is listed. The degrees of freedom is the number of experiments minus the number of coefficients.

### 16.11 *list* Parameter

List a specific parameter.

## 16.12 *list* Phase *<phase name>*

You must first specify the phase name. Then you can specify if you want the phase CONSTITUTION, DATA or some MODEL information.

### 16.12.1 list phase ... Constitution

List the constitution of the phase.

### 16.12.2 list phase ... Data

List the model and model parameter expressions.

### 16.12.3 list phase ... Model

List some model data for example if there is a disordered fraction set.

## 16.13 list Quit

You did not really want to list anyting.

#### 16.14 list Results

List the results of an equilibrium calculation. This is the most frequent list command. The listing will contain the current set of conditions, a table with global data, a table with component specific data and then a list of stable phases with amounts, compositions and possibly constitutions. It is possible to list also unstable phases.

There are 9 options for the formatting:

- 1 Output in mole fractions, phase constituents in value order (constituent with highest fraction first).
- 2 as 1 but include also the phase constitution (sublattices and their fractions) in value order.
- 3 as 1 with the phase composition in alphabetical order

- 4 Output in mass fractions, phase composition in value order.
- 5 as 4 with the phase composition in alphabetical order.
- 6 as 4 and also include the phase constitutions in value order.
- 7 Output all phases will with composition in mass fractions and in value order. Unstable phases will have a negative driving force.
- 8 Output all phases will with composition in mole fraction and constitution in alphabetic order. Unstable phases will have a negative driving force.
- 9 as 8 but in in value order.

For each phase the name, its status (S=suspended/D=dormant/E=entered/F=fix), moles (or mass), volume, number of formula units, atoms per formula units and driving force (in dimensionless units) is given on one line.

The moles of a phase is the number of formula unit multiplied with atoms per formula units. The gas phase and phases with interstitials and vacancies have a varying amount of moles of atoms per formula units. The composition of the phase can be in value order or alphabetical order.

### 16.15 list Short

There are 4 options: A/C/M/P

The A option lists a single line for each element, species and phase with some essential data.

The C option lists one line for each components.

The M option lists the models and constitution for all phases.

The P option lists one line for each stable phase and then one line for some of the remaining phases in decreasing order of stability.

#### 16.16 *list* State\_ Variables

Values of individual state variables like G, HM(LIQUID), X(LIQUID,CR) etc. can be listed. Terminate the command by an empty line. Note that the values of symbols and TP functions cannot be listed here, they are calculated by the CALCULATE SYMBOL or CALCULATE TP command.

The current values of parameter identifiers, see section 2.6.1 can be listed with the command, like TC(BCC) will give the calculated Curie temperature for BCC. A symbol like MQ&FE(FCC) will give the logarithm of the mobility of Fe in the FCC phase.

This command is superseeded by the SHOW command.

## 16.17 list Symbols

All state variable symbols listed but not their values, they are calculated by the CALCULATE SYMBOL command.

```
List of all state variable symbols
 No Special Name= expression ;
             R= 8.31451;
  1
  2
             RT = R*T;
  3
             T_C = T - 273.15;
  4
            CP = HM.T;
  5
            DCP= 1
          С
  6
         7X
            H298= HM;
```

In the "special" column the "D" means the symbol that is a "dot derivative" which is calculated only when explicitly specified, "C" means a numeric value that can be amended. The special 7X means a symbol that is evaluated only at equilibrium 7 which means you can refer to the value of this symbol calculated at the specified equilibrium in other equilibria. See also section 5.13.

## 16.18 *list* Tpfun Symbols

All or some TPFUN expressions listed. By giving \* all are listed, bu giving the g\* all TP functions starting with G are listed.

Note that all parameters are also TP functions, they can be listed by giving "\_\*" as name. The abbreviation "\_g\*" will list the function for all parameters with identifiers starting with G.

To obtain the values of TP functions use the **calculate TP** command.

# 17 Macro

By specifying a file name commands will be read from that file. The default extension is OCM. A macro file can open another macro file (max 5 levels). When a macro file finish with SET INTERACTIVE the calling macro file will continue or the user can continue interactively. See section 2.8.

When you start OC you can give a macro file name on the same line and the program will drictly start reading from this file.

With the popup window facility there are some special things. If you open the macro file with the popup window OC will save the directory where the macro file was found. If there are references to other files such as datbases or other macro files inside the macro and these file names are on the same line as the command **read tdb**./steel1 the file name must be

preceded by a "./", otherwise OC will try to open the file on its "working directory", see section 2.1.1.

# 18 Map

For phase diagram calculations. You must first set two axis with state variables which are already set as conditions.

If you give several MAP commands you can choose to erase or keep the previous results at each command.

During mapping each calculated equilibria is saved and for plotting any state variable can be used.

## 19 New

To remove all data so a new system can be entered. It is fragile.

# 20 Optimize

The model parameters selected by SET VARIABLE\_COEFF will be varied to obtain the best least square fit the experimental data provided.

Before this command you must have entered the thermodynamic descriptions of the phases with model parameters depending on optimizing coefficients and the experimental data. You must also set the range and weights of the experiments and which coefficients to be variable.

You must provid a maximum number of iterations allowed. If you give zero a "dry run" will be made with the current values of the optimizing coefficients. This can be useful to see that there are no problems calculating the equilibria.

Developing better assessment software is one of the main aspects of the OC software. There will be more options to this command soon.

# 21 Plot

Plot the result from a STEP or MAP calculation. A simple interface to GNUPLOT [GNUPLOT] has been implemented in OC. This generates a command file which is automatically plotted using GNUPLOT after the "render" command.

In OC you must first specify the state variable on the horizontal (x-axis) and vertical (y-axis) axis. Then you can give several of the options below, finish with RENDER or QUIT.

The simples way to generate a complex plot to be saved as PDF or PNG format is to first select the approriate axis and then set a few options like scaling, axis texts and text labels and plot on the screen. If you are not satisfied you can plot again (without changing the axis variables, if you change these all options you set will be cleared) and add or modify the options. When you are satisfied with the plot on the screen you plot a final time and set the GRAPHICS-FORMAT option and plot in the desired format on a file. Note that some formats may not be exactly identical as you see on the screen.

Default plotfile is "ocgnu.plt". This is the command file which will be executed by GNUPLOT. If GNUPLOT is correctly installed OC will start GNUPLOT and generate the graphics output when you RENDER the plot.

GNUPLOT is a very powerful graphics software, only a few of its facilities are available by OC. The gnuplot command file generated by OC can be edited to exploit additional facilities in GNUPLOT.

APPEND	LINE_WITH_POINTS	POSIION-OF-KEYS	TEXT
$AXIS_LABELS$	LOGSCALE	QUIT	TIE-LINES
FONT-AND-COLOR	MANIPULATE-LINES	RATIOS-XY	TITLE
GIBBS-TRIANGLE	OUTPUT-FILE	RENDER	
GRAPHICS-FORMAT	PAUSE-OPTIONS	SCALE-RANGES	

### A short summary:

- APPEND means overlay the current plot with another GNUPLOT file
- AXIS-LABELS you can specify the label on X or Y axis
- FONT-AND-COLOR you can select font and color of text
- GIBBS-TRIANGLE means an equilateral triangular diagram
- GRAPHICS-FORMAT to select the GNUPLOT output device (PS, PDF, PNG etc)
- LINE-WITH-POINTS means a symbol at every calculated equilibrium
- LOGSCALE you can specify that X or Y axis is logaritmic
- MANIPULATE-LINES is to select line identification
- OUTPUT-FILE the GNUPLOT file is saved on this file (default ocgnu.plt)
- PAUSE-OPTION to select how GNUPLOT should behave after plotting
- POSITION of the identification labels for the curves
- QUIT no plot generated
- RATIOS-XY will change the relative length of X and Y axis

- RENDER finally plot
- SCALE-RANGES for X and Y axis you can specify min and max value plotted
- TEXT you can place a text inside the plot
- TIE-LINES if you have tie-lines in the plane you can plot some of them
- TITLE the heading of the plot

## 21.1 plot xaxis yaxis Append

A GNUPLOT file prevously generated by OC or edited manually can be specified to be overlayed on the current plot.

## 21.2 plot xaxis yaxis Axis\_Labels

You specify for the X or Y axis the axis labels. By default the state variable or symbol plotted will be used as label.

For X or Y axis?

Specify the axis for which you want to enter the label

Axis label:

The default label is given in the question.

# 21.3 plot xaxis yaxis font-and-color

Not implemented yet. Intended to select font and color of all text. Some GNUPLOT terminals may not support certain fonts.

# 21.4 plot xaxis yaxis Gibbs triangle

Gibbs triangle plots can only be used for isothermal sections. A trial implementation is available which can generate equiaxial triangular isothermal diagrams. If you already set this the default is to plot on a square.

# 21.5 plot xaxis yaxis Graphics format

The GNUPLOT terminals entered in section 10.8 can be used. For other formats than SCREEN you can also specify an output file which will be generated in the specified format.

Graphics format index:

The default terminal indices are 1 SCREEN

- 2 PS (Postscript)
- 3 PDF (Adobe Portrable Document Format)
- 4 GIF
- 5 PNG

You can change these or enter more graphics formats by the **enter gnuplot** command. 10.8.

If SCREEN is not selected the GNUPLOT program with generate a file with the plot and you can specify the name of this file. It will have the appropriate extention depending on the format.

Plot file:

In addition to the GNUPLOT command file the graphics a file with the specified format will be generated.

## 21.6 plot xaxis yaxis Line-with-points

Not implemente yet

## 21.7 plot xaxis yaxis Logscale

You can set logarithimic scale on X or Y axis (or both).

# 21.8 plot xaxis yaxis Manipulate lines

This is not implemented. It is intended to allow specification of the color of the curves in the plot.

# 21.9 plot xaxis yaxis Output file

By default plotting will generate a ocgnu.plt file for GNUPLOT. You can specify other name here. If you plot on other terminals than SCREEN there will be an additional file with extension ".ps" for Postscript, ".pdf" for Adobe PDF or ".gif" for GIF format.

# 21.10 plot xaxis yaxis Pause option

When you plot on the screen the last command on the file to GNUPLOT is "pause mouse". You can change this with this command.

## 21.11 plot xaxis yaxis Position of keys

The identification (labels) of the curves in the plot can be positioned with this command. See the GNUPLOT manual [GNUPLOT] for information.

## 21.12 plot xaxis yaxis Quit

No plot generated.

## 21.13 plot xaxis yaxis Ratios XY

The relative ratios of the X and Y axis can be specied.

## 21.14 plot xaxis yaxis Render

Finally plot using all the option set.

## 21.15 plot xaxis yaxis Scale\_Range

You specify for the X or Y axis the minimum and maximum range. The automatic (default) scaling range can always be restored.

# 21.16 plot xaxis yaxis Text

This is a facility to add a text to a plot at an arbitrary position.

## 21.16.1 plot xaxis yaxis text Modify existing text?:

If there is already some text items you must first answer if you wants modify an already existing one. If so all the texts are listed and you can select which one you wants to modify.

### 21.16.2 plot xaxis yaxis text Which text index?:

You must provide the index of the text to change.

For a new or changed text you must give:

### 21.16.3 plot xaxis yaxis text X position

The X coordinate of the text (in the plot scale)

#### 21.16.4 plot xaxis yaxis text Y position

The Y coordinate of the text (in the plot scale)

### 21.16.5 plot xaxis yaxis text Fontscale

A relative size factor, default is 1. The size of the text will be scaled accordingly.

#### 21.16.6 plot xaxis yaxis text Angle (degrees)

The text will be written with the specified angle. Zero means horisontally, negative valus slopes downward, positive upwards. An ange of 180 means the text will be upside down.

#### 21.16.7 plot xaxis yaxis text Do you want to calculate the equilibrium?

If it is a phase diagram that is plotted you can select to calculate an equilibrium at the specified coordinates. The names of the stable phases will be proposed as text.

### 21.16.8 plot xaxis yaxis text Text:

The text to be added to the plot. The text will start at the coordinates given. On Postscript and PDF a greek character can be given as "/Symbol m" for  $\mu$ .

## 21.17 plot xaxis yaxis Tie-line

Tie-lines in isothermal ternary phase diagram can be plotted.

Tie-line plot increment?

The increment is related to the actual equilibria calculated. 0 means no tie-lines plotted, 3 means to plot a tie-line at every 3rd calculated equilibria and so on.

# 21.18 plot xaxis yaxis Title

Title on top of the figure. The conditions will always be there also.

# 22 Quit

Terminate the OC software in English, have a nice day.

## 23 Read

It is possible to read a (non-encrypted) TDB file but it should be not too different from what is normally generated by the LIST\_DATA command in TC.

```
DIRECT PDB UNFORMATTED QUIT TDB
```

#### 23.1 read Direct

In the future it will be possible to save results on a random access (DIRECT) file.

## 23.2 read Quit

You did not really want to read anything.

### $23.3 \quad read \text{ PDB}$

A PDB file (with extension PDB) should be specified. The file should be un the Portable phase dependent Data Base format.

### $23.4 \quad read \text{ TDB}$

A TDB file (with extension TDB) should be specified. The TDB file must not deviate very much from the standard output from Thermo-Calc.

File name:

If you do not use the popup window for opening files you must specify the database file name. The file must be on the working directory (where you started the OC program, see section 2.1.1) or you must provide the path.

After opening the file the program will list the elements and ask: Select elements /all/:

If you give RETURN the data for all elements will be read. If you specify one or more elements data for those will be read. You cannot select the phases here but later you can suspend those you are not interested in.

### 23.5 read Unformatted

For use to read a file created with a SAVE UNFORMATTED command. It may not always work as the data structure is still changing.

## 24 Save

There are several forms of save, three forms write a text file that can be read and modified with a normal editor. Two forms are unformatted, either on a sequential file or a direct (random access) file.

DIRECT SOLGAS UNFORMATTED QUIT TDB PDB

#### 24.1 save Direct

It will eventually be possible to save the result of STEP and MAP commands on a random access file for later processing.

## 24.2 save Quit

You did not want to save anything.

### 24.3 save PDB

Saves current set of model parameters and functions on a file in the Portable phase dependant Data Base format.

## 24.4 save TDB

Saves current set of model parameters and functions on a file in TDB format. Same as the command list data tdb.

### 24.5 save SOLGAS

Saves current set of model parameters and functions on a file in a format that (hopefully) can be read by the FactSage software.

### 24.6 save Unformatted

With this command you can save the current status of the calculations on a file and then resume the calculations by reading this file. Note that the Fortran unformatted files may not be portable, they depend on the compiler, the operating system and the hardware.

# 25 Select

There are a few things that can be selected, most important which equilibrium the following commands will operate on.

## 25.1 select Equilibrium

As you can enter several equilibria with different conditions this command allows him to select the current equilibria.

## 25.2 select Graphics

Only GNUPLOT citegruplot available.

## 25.3 select Language

Only English implemented (except a few French exclamations).

### 25.4 select Minimizer

Only Hillert's algorithm implemented in matsmin [15Sun2] available.

# 25.5 select Optimizer

The LMDIF [LMDIF] least square fitting software is the only one implemented.

## **26** Set

Many things can be set. Things to be "set" and "amended" sometimes overlap.

ADVANCED	$FIXED\_COEFF$	OPTIMIZING_COND	STATUS
AS_START_EQUILIB	INPUT_AMOUNTS	PHASE	$T_AND_P$
AXIS	INTERACTIVE	QUIT	UNITS
BIT	LEVEL	RANGE_EXP_EQUIL	VARIABLE_COEFF
CONDITION	LOG_FILE	REFERENCE_STATE	VERBOSE
ECHO	NUMERIC_OPTIONS	$SCALED\_COEFF$	WEIGHT

### 26.1 set Advanced

A few options implemented

### 26.1.1 set advanced equilibrium transfer

This is only for experts who know what they are doing.

#### 26.1.2 set advanced extra property

No implemented yet.

### 26.1.3 set advanced global-min-onoff

Turn on or off using global minimization.

### 26.1.4 set advanced grid\_density

At present the grid density cannot be fine tuned. For some phases it is fixed for some you can select a more or less dense grid.

#### 26.1.5 set advanced map-special

Not implemented yet.

### 26.1.6 set advanced popup-window

Any other answer then Y will turn them off. By answering Y you turn on popup windows for opening files (the default).

#### 26.1.7 set advanced quit

You did not want to set anything advanced.

### 26.1.8 set advanced small-grid-onoff

A particularly small grid for the global minimization is selected or not.

### 26.1.9 set advanced working-directory

The name of the working directory (where OC was started) is listed. It cannot be changed at present. It is related to the popup windows for opening files, see section 2.1.1.

## 26.2 set As start equilibrium

The current equilibrium will be copied to the list of start equilibria for STEP and MAP commands.

#### 26.3 set Axis

A condition can be set as an axis variable with a low and high limit and a maximum increment. With 2 or more axis you will calculate a phase diagram, i.e. lines where the set of stable phases changes.

With one axis you calculate the set of stable phases and their properties while changing the axis variable.

#### 26.4 *set* Bit

Many records have status words where the bits are used to signify different things. An advanced user can set these bits for the global, equilibrium and phase records, but only if you know what it means.

- The global record bits are listed below. Most of them are set or reset automatically by the software or by other commands.
  - 0 you are a beginner (set by default)
  - 1 you are experienced
  - 2 you are an expert
  - 3 gridminimizer must not be used
  - 4 gridminimizer must not merge comp.sets. (set by default)
  - 5 there are no data (cleread automatically)
  - 6 there are no phases (cleared automatically)
  - 7 comp.sets must not be created automatically
  - 8 comp.sets must not be deleted automatically
  - 9 data has changed since last save (set automtically)
  - 10 means verbose is on (not implemented)

- 11 means verbose is permanently on (not implemented)
- 12 means be silent (supress warnings)
- 13 no cleanup after an equilibrium calculation
- 14 use denser grid in grid minimizer (see also SET ADVANCED)
- 15 calculations in parallel is not allowed
- 16 no global test at node point during STEP/MAP
- 17 the components are not the elements
- 18 global test of equilibrium AFTER calculation
- 19 use old (less dense) grid minimizer
- 20 do not recalculate if global test AFTER fails
- 21 use old MAP algorithm
- -22-31 not yet used
- The EQUILIBRIUM record bits are listed below
  - 0 No threads allowed (no parallel calculation)
  - 1 No global minimization allowed for this equilibrium
  - 2 No equilibrium has been calculated (there are no results)
  - 3 Conditions and results not consistent
  - 4 Last equilibrium calculation failed
  - 5 No automatic generation of composition sets
  - 6 Equilibrium tested by grid minimizer
  - 7 Current results are from a grid minimization
- To change the phase status word use SET PHASE ... bit

### 26.5 set Condition

A condition is a value assigned to a state variable or an expression of state variables. By setting the status of a phase to fix you have also set a condition.

Two cases of expressions can be used as conditions, for example a relation between mole fraction like

### set condition $x(liq,o)-x(c1_mo2,o)=0$

means that the oxygen content in liquid and c1\_mo2 phases should be the same. That is useful to calculate the congruent melting of c1\_mo2.

Another case is if the total amount if some components has a relation, for example:

## set condition n(u)+n(zr)=1

means that the total number of moles of the components U and Zr should be unity.

#### 26.6 set Echo

This is useful command in macro files or when demonstrating the program.

### 26.7 set Fixed coefficient

An optimizing coefficient is assigned a fixed value.

### 26.8 set Input Amounts

Output for equilibrium:

This allows you to specify a system by giving a redundant amount of various species in the system. The software will transform this to conditions on the amounts of the components. An example:

```
--->OC5:read tdb cho-gas
--->OC5:set input n(c1o2)=10
--->OC5:set input n(c1h4)=5
--->OC5:1 c
Conditions for equilibrium: 1, DEFAULT_EQUILIBRIUM
1:N(C)=45, 2:N(O)=80, 3:N(H)=30
Degrees of freedom are 2
```

The amounts of the species has been split on the components. Setting input amounts is just another way to set these directly. If we set a T and P we can calculate the equilibrium fraction of all the species.

```
--->0C5:set c t=1000 p=1e5
--->0C5:1 c
Conditions for equilibrium:
                              1, DEFAULT_EQUILIBRIUM
  1:N(C)=45, 2:N(0)=80, 3:N(H)=30, 4:T=1000, 5:P=100000
 Degrees of freedom are
--->0C5:c e
 3Y Constitution of metastable phases set
                          1.56E-02 s and
                                                0 clockcycles, T= 1000.00
Gridmin:
              85 points
Phase change: its/add/remove:
                                   5
                                             0
                                       11
Phase change: its/add/remove:
                                             0
                                  12
                                       12
Phase change: its/add/remove:
                                  17
                                        0
                                            12
Phase change: its/add/remove:
                                  53
                                        0
                                            11
Equilibrium calculation
                          79 its,
                                    7.8125E-02 s and
                                                           93 clockcycles
--->0C5:1
LIST what? /RESULTS/:
Results output mode: /1/:
```

1, DEFAULT\_EQUILIBRIUM

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```
Conditions .....:
  1:N(C)=45, 2:N(O)=80, 3:N(H)=30, 4:T=1000, 5:P=100000
Degrees of freedom are
Some global data, reference state SER ............
T=
    1000.00 K (
                 726.85 C), P= 1.0000E+05 Pa, V= 4.9872E+00 m3
N=
    1.5500E+02 moles, B= 1.8507E+03 g, RT= 8.3145E+03 J/mol
GS= -2.80411E+07 J, GS/N=-1.8091E+05 J/mol, HS=-1.2914E+07 J, SS= 1.513E+04 J/K
Some data for components .....:
                           Mole-fr Chem.pot/RT Activities Ref.state
Component name
                Moles
С
                4.5000E+01 0.29032 -3.7354E+00 2.3863E-02 SER (default)
Η
                3.0000E+01 0.19355 -9.8098E+00 5.4910E-05 SER (default)
0
                8.0000E+01 0.51613 -3.6377E+01 1.5911E-16 SER (default)
Some data for phases .....:
Name
                  Status Moles
                                            Form.Units Cmp/FU dGm/RT Comp:
                                   Volume
GAS..... E 1.550E+02 4.99E+00 6.00E+01
                                                         2.58 0.00E+00 X:
       5.16129E-01 C
                          2.90323E-01 H
                                            1.93548E-01
Constitution: There are
                          73 constituents:
C102
             4.54395E-01 C2H3
                                     8.67456E-17 C4H10_1
                                                              2.73242E-23
C101
             2.95682E-01 C3H4_2
                                     3.04922E-17
                                                 C4H10_2
                                                             1.38822E-23
H201
             1.29270E-01 C3H8
                                     2.73523E-17 C4H2
                                                             8.16657E-24
H2
             1.20501E-01 C3H6O1
                                     1.94895E-17 H102
                                                             4.37267E-24
             1.52786E-04 C3H4_1
                                     8.18695E-18 C4H6_5
C1H4
                                                             1.44915E-24
C1H2O2_CIS
             4.04887E-08 C1H3O1_CH3O 3.87833E-18 C4H8
                                                              1.04297E-25
             2.01368E-08 C2H4O1_OXIRA 1.64221E-19 C2H1
C1H2O1
                                                             7.79712E-26
C1H2O2_TRANS 5.82767E-09 C1H2
                                     3.98656E-20 C4H8_4
                                                             6.39692E-26
Η
             7.88542E-10 H202
                                     3.27068E-20 C6H6O1
                                                              3.00598E-26
C1H4O1
             1.27636E-10
                                     1.46838E-20 C1H1
                                                              1.81712E-27
 C2H4
             1.05140E-10 C2H6O2
                                     1.19305E-20 C3H1
                                                              1.68523E-28
 C2H6
             3.44726E-11 02
                                     8.71930E-21 C4H4_1_3
                                                             7.73762E-29
C1H3
                                     5.73533E-21 C1H2O2_DIOXI 4.04963E-30
             1.83302E-11 C4H6_2
C1H1O1
             7.24719E-12 C201
                                     1.72590E-21
                                                 C4H1
                                                              1.00000E-30
                                                 C2H4O2_DIOXE 1.00000E-30
C2H4O1_ACETA 2.00054E-12 C4H8_5
                                     9.38081E-22
H101
             1.86354E-12 C4H8_3
                                     5.91323E-22 C4
                                                              1.00000E-30
 C2H2
             1.82837E-12 C4H8_1
                                     4.75317E-22 C2H4O3_123TR 1.00000E-30
                                     4.17043E-22 C2H4O3_124TR 1.00000E-30
C1H102
             1.57298E-12 C4H8_2
C2H4O2_ACETI 7.65642E-13
                         C2H2O1
                                     1.47405E-22 C2
                                                              1.00000E-30
C1H3O1_CH2OH 1.64978E-15 C4H6_4
                                     8.47392E-23 C60
                                                              1.00000E-30
             1.11079E-15 C6H6
                                     8.21607E-23 C3
                                                             1.00000E-30
C302
             7.21243E-16 C4H4
                                     5.46648E-23 C5
C3H6_2
                                                              1.00000E-30
 СЗН6
             7.13743E-16 C4H6_1
                                                             1.00000E-30
                                     5.05773E-23 03
 C2H601
             6.22811E-16 C4H6_3
                                     2.87604E-23
 C2H5
             4.72671E-16 C4H10_1
                                     2.73242E-23
```

The calculation shows that mixing 10 moles of  $CO_2$  with 5 moles of  $CH_4$  at 1000 K and 1 bar gives a gas with 45%  $CO_2$ , 30% CO, 13%  $H_2O$  and the rest  $H_2$ 

#### 26.9 set Interactive

The last command on a macro file. Gives command back to the keyboard of the user, or to the calling macro file. Without this the program will just terminate when the macro is finished.

### 26.10 set Level

I am no longer sure what this should do and if it is needed ...

### 26.11 set Log\_File

A useful command to save all interactive input while running OC. The log file can easily be transformed to a macro file. All bug reports should be accompanied by a log file which reproduces the bug.

## 26.12 set Numeric\_Options

The default number of iterations and accuracy can be specified. Default values are 500 and  $10^{-6}$ .

# 26.13 set Optimizing conditions

A few variables used to guide the optimization of model parameters can be set.

# 26.14 set Phase <phase name>

You must specify a phase name. Some phase specific things can be set, also for the model.

### 26.14.1 set phase ... Amount

You can specify the amount of the phase which is used as initial value for an equilibrium calculation.

#### 26.14.2 set phase ... Bits

Some of the models and use of data storage depend on the bits of the phase. Most of them are set automatically by the software and other commands like AMEND PHASE. Changing them with this command will not have the expected effect and may cause the program to fail.

The bits that can be changed are:

- EXTRA\_DENSE\_GRID makes it possible to have a larger number of gridpoints calculated by the gridminimizer for the specified phase.
- NO\_AUTO\_COMP\_SET. This makes it possible to prevent that the specific phase has automatic composition set created during calculations.
- OUIT, do not set any more bits.

### 26.14.3 set phase ... Constitution

This is the same as **amend phase constitution**.

### 26.14.4 set phase ... Default\_constitu

Same as amend phase default\_constit.

#### 26.14.5 set phase ... Quit

You did not want to set anything for the phase.

#### 26.14.6 set phase ... Status

Use the SET STATUS PHASE command to set the status of one or several phases. The different status are explained for that command.

### 26.15 *set* Quit

You did not really want to set anything.

# 26.16 set Range of experimental equilibria

For an assessment several consequtive equilibria with experimental data must be entered. This command specifies the first and last of those equilibria. It possible to add more equilibria later one by one (not yet though).

The equilibria are assigned the weight one by default. The weight can be changed with the SET WEIGHT command. The weight zero means the equilibrium is not calculated.

#### 26.17 set Reference\_State

By default the reference state for the components is SER (Stable Element Reference) which is the stable state of the element at 298.15 K and 1 bar. (NOTE: in principle SER is defined by the database but today almost all databases have SER as reference state.)

For each component (also for other components than the elements) you can specify a phase at a given temperature and pressure as reference state. The phase must exist for the component as pure. Instead of a fixed T you can give a \*, indicating current T, if you calculates at different values of T. Example:

#### set reference O gas \* 1e5

Note that state variables like the chemical potential, MU(O), will refer to the user defined reference state. To obtain the value for the SER state you can use the suffix S, i.e. MUS(O) will give the chemical potential referred to SER.

IMPORTANT NOTE: the value of integral properties like Gibbs energy, G, enthalpy, H, etc. will also be affected by the change of the reference state of an element. If all elements have the same phase as reference state the value of the enthalpy obtained by H for that phase will be the enthalpy of mixing.

In order to have use SER as reference state use a suffix S. The enthalpy relative to SER is HS independent of any reference state set for the elements.

### 26.18 set Scaled coefficient

A coefficient for optimization can be specified with a start value, scaling factor and a minimum and maximum value. The *set* VARIABLE command sets the scaling factor equal to the start value and have no min or max values.

#### 26.19 set Status

The status of elements, constituents, species or phases can be changed. Only phases are implemented.

#### 26.19.1 set status Constituent

A constituent of a phase can be suspended. Not yet implemented.

#### 26.19.2 set status Element

An element can be ENTERED or SUSPENDED. If an element is suspended all species with this element is automatically suspended. Not yet implemented.

#### 26.19.3 set status Phases

A phase can have one of 4 different status

- ENTERED, this is the default. The phase will be stable if that would give the most stable state for the current conditions. The user can give a tentative amount.
- SUSPENDED, the phase will not be included in any calculations.
- DORMANT, the phase will be included in the calculations but will not be allowed to become stable even if that would give the most stable equilibrium. In such a case the phase will have a positive driving force.
- FIXED means that it is a condition that the phase is stable with the specified amount. Note that for solution phases the composition is not known.

You can use a list of phase names or a wildcard for the phase name and the must give an equal sign, "=", before the new status. You can also use the special "\*S" for all suspended phase, "\*D" for all dormant phases.

Changing the phase status does not affect anything except the phase itself. For a single phase you can use SET PHASE ... STATUS <status>.

Setting a stable phase as dormant or suspended and calculate the equilibrium will give you a metastable equilibrium.

Setting a phase status as FIXED means it is a condition that this phase should be stable. Setting the liquid fix with the amount zero is a quick way to calculate the melting temperature of a system if there is no condition on the T.

#### 26.19.4 set status Species

A species can be ENTERED or SUSPENDED. If a species is suspended all phases that have this as single constituent in a sublattice will be automatically suspended. Not yet implemented.

### 26.20 set $T_and_P$

Local values of T and P can be set. These are not conditions but are used for commands like **CALCULATE PHASE** ....

### 26.21 set Units

For each property the unit can be specified like Kelvin, Farenheit or Celsius for temperature. Not implemented yet.

### 26.22 set Variable coefficient

A coefficient Aij for optimization is assigned a start value or its current value to be optimized against selected experimental data.

### 26.23 set Verbose

Not implemented yet.

## 26.24 set Weight

Intended for assessments. A weight is zero or a positive value. Equilibria with weight zero will be ignored in an optimization.

You can specify the current equilibrium or give an abbreviation that will set the weight of all equilibria with a name for which the abbreviation fits. Or you can give a range of equilibria by giving two numbers separated by a hyphen like 63-106.

If an abbreviation or a range is given the software will list how many equilibra that had the weight set to the new value.

# 27 Show

This command shows a value of a property, the property can be a state variable like T, G etc or a user detfined symbol containing several state variable or a model parameter identifier (which must always have a phase specification) like the Curie temperature. The state variables can contain wildcards like X(FCC,\*) means all mole fractions of the FCC phase. Several properties can be specified on the same line, separated by a space character.

## 27.1 property:

One or more properties can be specified like

show t g tc(bcc) mu(cr)

# 28 Step

Requires that a single axis is set. If a second step command is given you have the choice of deleting or keeping the previous results.

## 28.1 step Conditional

A specified symbol is evaluated at each step. Can be used for Scheil-Gulliver solidification simulation when implemented.

## 28.2 step Normal

Calculates equilibria from the low axis limit to the high at each increment. The exact axis value for any phase changes is calculated.

## 28.3 step Quit

You did not want to step.

# 28.4 step Separate

Calculates each phase separately. It calculates equilibria for each phase separately. It can be used to calculate Gibbs energy curves.

# 29 Summary

That's all and I hope enough (when all is implemented). Have fun and report all errors or problems providing a macro file and the necessary data.

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