

THERMOCALC

everything you wanted to know but were too afraid to ask

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Script File Documentation

The information below pertains to using scripts in THERMOCALC (for scripts in drawpd, see the drawpd documentation). This list is an updated version of the list given in the course CD documentation originally compiled by Roger Powell and updated now by Richard White.

Clicking a script below will take you to a more indepth explanation of what the script does and what commands can follow it.

Overview

Scripting allows simple customisation of the running of THERMOCALC. This is a good idea because each time you run the program you would otherwise be met with a barrage of questions many of which you might know in advance are not relevant or that can be set ahead of time. It is usually possible to run THERMOCALC without using scripts, but using them can make your life much easier. In addition some facilities are only available via scripts.

There are two places that scripts are read by THERMOCALC :

- 1. in a separate file called tc-prefs.txt: these scripts control global aspects of running THERMOCALC. Scripts here will affect all runs of THERMOCALC with different datafiles.
- 2. in the scriptfile (or the second section of a datafile in older THERMOCALC versions): these local scripts control the way THERMOCALC runs that particular axfile.

Each of the controls which can be set by scripts are set to a default in THERMOCALC. All that happens with the provision of scripts in tc-prefs or in a datafile is that the default is overridden. Each time it is run, THERMOCALC reads these scripts, and these override the defaults; THERMOCALC does not store or remember previously read scripts.

The basic script format involves a single line of file, starting with a "keyword". If you don't get the keyword right (ie if THERMOCALC doesn't recognise it), then the line

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is ignored and (generally) there will be a line of output at the beginning of a run that this has happened "script unrecognised" (but THERMOCALC won't bomb).

The simplest form of script involves a "keyword" followed by one of "yes", "no" or "ask". "Yes" corresponds to some action which will definitely be undertaken, and "no" corresponds to the action definitely not being undertaken. With "ask", you will be prompted by THERMOCALC, while the program is running, to determine if the action should be undertaken. In some cases "no" gets substituted for "ask" with no prompt allowing the meaning of "ask" to be clarified. If one of "yes", "no" or "ask" is omitted, then "yes" is assumed.

More complex scripts allow additional information to be provided to THERMOCALC. Remember that if THERMOCALC doesn't understand what you have tried to do with a script, there is no problem, but that the intended action or inaction will not occur. This will normally be obvious in the way THERMOCALC runs! As each script is a single line of datafile, one script which doesn't work will not affect succeeding scripts. (In the following, the forms of scripts are in quotes—don't include the quotes when you use the scripts)

Below is a list of scripts as at Dec 2007 for THERMOCALC 3.30. While many scripts here work in earlier versions, not all of them wil. The links in each script will take you to a more in-depth explanation of what the script does, what script suffixes can be used and how to use each script. The scripts are separated into two groups, being those for the preferences file and those for the script file. The scripts serve a range of purposes including telling THERMOCALC what files to use, setting what types of calculations to run, what types of output to produce, what information to include in that output and controlling things such as minerals to use, P-T windows etc. A few scripts that have been recently superceded or no longer work in THERMOCALC 3.30 but are relevant to recent versions such as 3.23-3.26 are included

For calculating particular types of diagrams, ready-made script files with the set of scripts required are available for download in the datafiles page.

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Scripting in the preferences file

The most useful scripts which can be used in th prefs are:

Script Name	Short Description
dataset	specifying which version of the internally consistent dataset to use (currently 55, meaning that tc-

	ds55.txt is used)
<u>scriptfile</u>	specifying which script file suffix to use (replaces datasuffix)
<u>setpagewidth</u>	allows the width of the calculation window to be set (80 for the pc black window; whatever size is used for the Terminal window on the Mac)
<u>dontwrap</u>	whether line wrap occurs in screen output (for specified pagewidth)
<u>calcmode</u>	specifying a calculation mode for all runs
strictsyntax	better ax code checking, but involves a strict naming convention for variables

Useful preference file scripts

Script Name	Short Description
datasuffix	specifying a datafile suffix
pdatasuffix	specifying a suffix on tc <dataset> eg ds55</dataset>

main pre 3.30 scripts (equivalent to some of the above table

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Scripting in mode 1 datafiles

Scripts for phase diagram calculations that may appear in script files. Clicking on a script takes you to a more indepth explanation of the script and what suffixes can/must follow it.

Defining the a-x file to use

Script Name	Short Description
<u>axfile</u>	defining the a-x file to use

Scripts relating to calculation of P-T window

Script Name	Short Description
	defining the default P

setdefPwindow	window for calculations
<u>setdefTwindow</u>	defining the default T window for calculations
<u>setPwindow</u>	defining the P range for calculations
setTwindow	>defining the T range for calculations
<u>calctatp</u>	whether to calculate T at given P for PT-lines
moreprec	specifying more output precision (now standard in version 3.30)
calcsdnle	whether to perform uncertainty propagation on calculations

Calculating P-T window

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Scripts related to phases involved in calculations

Script Name	Short Description
omit or ignore	specifying phases to ignore this run
fluidpresent	whether H2O or CO2 in the datafile means fluid present
fluidexcess	whether fluid is "in excess"
setexcess	defining other "in excess" phases
exbuff	allows external buffering calculations involving H2O-CO2 fluids
<u>seta</u>	whether to set activities in calculations
<u>setmu</u>	whether to set chemical potentials in calculations
usecalcq	whether to pre-calculate the state of order in order-disorder phases
samecoding	whether to line up output for minerals with the same a-x code for easy inspection

Scripts related to phases involved in calculations

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Scripts related to calculating pseudosections and setting bulk rock composition

Script Name	Short Description
pseudosection	whether to generate pseudosection information
setbulk	defining bulk composition(s) for pseudosections
<u>setmodeiso</u>	allows the setting of mode isopleths
zeromodeiso	allows the setting of zero mode isopleths
<u>setiso</u>	whether to calculate composition isopleths in calculations
calcg	Gibbs energy of equilibrium calculated
dogmin	allows Gibbs energy minimization
<u>rbi</u>	allows a matrix of calculated mineral modes and mineral compositions to be used as the bulk rock composition
calcah2o	whether to calculate the activity of water
calcmu	whether to calculate the chemical potential of each component

Scripts related to calculating pseudosections and setting bulk rock compositions

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Output for drawpd

Script Name	Short Description
drawpd	calculated coordinates into the file called tc-blah-dr, where blah is the name given in the scriptfile script (or prompted)

Scripts related to output for drawpd

Related to calculating projections

Script Name	Short Description
project	whether to project phases onto a compatibility diagram
<u>setproject</u>	defining the compatibility diagram (projection plane)
<u>projcomp</u>	whether to project bulk comps onto the compatibility diagram

Scripts related to calculating projections

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Optional output in the log file

Script Name	Short Description
printbulkinfo	whether to include a bulk composition matrix in the logfile
printguessform	whether to include the output mineral compositions in starting guess form in the logfile (superceded by printxyz)
<u>printxyz</u>	whether to include the output mineral compositions in starting guess form in the logfile
incthermo	whether to include tables of thermo data in the logfile
incax	whether to include the a- x relationships in the logfile
incnler	whether to include the set of reactions in the logfile
<u>lsqds</u>	allows activities of end- members rather than mineral comps to be output for calculations

Optional output in the log file

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Scripts to produce Mathematica output (mainly for wizards)

Script Name	Short Description
<u>smath</u>	calculated PT coordinates into 'other' file
xinsmath	calculated PT and x coordinates into 'other' file
tabsmath	calculated PT coordinates into 'other' file in tab-separated form
<u>cmath</u>	compatibility diagram coordinates into 'other' file
txmath	Tx/Px pseudosection coordinates into 'other' file

Scripts to produce Mathematica output

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Scripts for avPT (mode 2)

Script Name	Short Description
<u>defsdaparams</u>	for setting activity uncertainty defaults
<u>usedefsda</u>	using the defaults
<u>justrelu</u>	just relative uncertainties (as in the thermobarometry paper)
setrockname	allows the results to be labelled with the rock name
diagnostics	to always see the diagnostics

Scripts for avPT (mode 2) calculations

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Scripts for calculations not involving (solid) solutions (mode 3)

Script Name	Short Description	
<u>setmu</u>	allows you to set the chemical potentials of components	
seta / setlna / setloga	allows you to set activities of components	

Scripts for calculations not ivolving (solid)

solution (mode 3)

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Scripts Explanations

Below is a list of usable scripts as at Dec 2007 for THERMOCALC 3.30. While most scripts here are usable in earlier versions, not all of them will be (so using THERMOCALC 3.30 is recommended). The links in each script will take you to a more in-depth explanation of what the script does, what script suffixes can be used and how to use each script. The scripts are separated into two groups, being those for the preferences file and those for the script file. The scripts serve a range of purposes including telling THERMOCALC what files to use, setting what types of calculations to run, what types of output to produce, what information to include in that output and controlling things such as minerals to use, P-T windows etc.

Download a pdf file of script explanations

More information about the <u>file structure for input and output files.</u>

Scripting in the preferences file

dataset (for version 3.30 onward)

The file with the prefix/root tc-ds is the internally consistent dataset (eg tc-ds55) that is to be used in the calculations. In most cases, this does not change very often (eg tc-ds55 is from Nov. 2003). For calculations, the dataset to be used is specified in the tc-prefs file. Forms

Form	What it does	
"dataset blah"	where "blah" is the text that follows the "tcds" (eg "dataset 55" to use tc-ds55)	

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scriptfile (for version 3.30 onward)

This script is available if you are spending a lot of time working with just one scriptfile; it allows you to specify this file in tc-prefs, via the suffix to the root of the datafile name. Having done this, it means that you can only work with this file (until you change the script, or remove it). This root is either the default, 'tc-', or whatever is specified in the script datafilename. If "scriptfile" is not present in tc-prefs, then the suffix to 'tc-<name>-o' (or whatever is specified in the script datafilename) is

prompted for when you run THERMOCALC. Form:

Form	What is does	
"scriptfile blah"	scriptfile used is tc-blah (Win: tc-blah.txt)	

Default: datafile name suffix will be prompted for

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setpagewidth

Allows the width of the calculation window to be set (80 for the pc black window; whatever size is used for the Terminal window on the Mac)

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dontwrap

Screen output on smaller screens (or longer lines of output) can look messy if a line is wrapped onto the following one. This script allows user control over this. With "outdialog yes", a # is placed at the end of a line which would otherwise wrap onto the following line. The full line of course does appear in the output file. Use in conjunction with "dontaskadjust", to control the width of window on which the wrapping is not done. Forms:

Form	What it does	
"dontwrap yes"	don't wrap output on screen	
"dontwrap"	don't wrap output on screen	
"dontwrap no"	allow wrap of output on screen	

dontwrap options

Default: "dontwrap yes".

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calcmode

If you are only doing one type of calculation, say only average PT calculations, then you can specify this with this script (and save yourself always replying in this way to a prompt). Form:

Form	What is does	
"calcmode -1"	calculation mode prompted for (this is the equivalent of the script not occuring in th prefs)	

"calcmode 0"	data table mode (this would not be a usual setting)
"calcmode 1"	phase diagram calculations involving phases which are (solid) solutions
"calcmode 2"	rock calculations (ie average PT calculations)
"calcmode 3"	calculations involving all the reactions between a set of end-members (for phase diagrams not involving (solid) solutions and debugging average PT calculations)

Default: calculation mode prompted for

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strictsyntax

allows better ax code checking, but requires a strict naming convention for all variables.

Form	What it does
"strictsyntax yes"	strict syntax checked for
"strictsyntax"	as above
"strictsyntax no"	allows a less rigid ax coding

options for strictsyntax script

Default: strictsyntax yes

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datasuffix (for pre version 3.30)

This script is available if you are spending a lot of time working with just one datafile; it allows you to specify this file in th prefs, via the suffix to the root of the datafile name. Having done this, it means that you can only work with this file (until you change the script, or remove it). This root is either the default, 'th d', or 'tcd' or whatever is specified in the script datafilename. If "datasuffix" is not present in th prefs, then the suffix to 'th d' (or whatever is specified in the script datafilename) is prompted for when you run THERMOCALC. Form:

From	What it does	
"datasuffix	datafile used is th dblah	
blah"	(Win: thblah.txt)	

Default: datafile name suffix will be prompted for

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pdatasuffix (pre version 3.30)

The pdata file contain the thermodynamic data which THERMOCALC uses for all its calculations. As the pdata files are updated, the results of calculations change (if usually only slightly). Sometimes it is useful to keep older pdata files, so that original calculations can be reproduced. This script allows several generations of pdata files to be kept together, using different suffices on the pdata filenames, and the choice of which ones are used being controlled by a tcprefs script. Forms:

From	What it does
"pdatasuffix blah"	pdata file will be expected to have a name like tc-ds55 (dataset 5.5)

Default: files are named tc-ds55

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Scripting in mode 1 datafiles

axfile

This script sets the a-x file to be used in the calculations. A-x files are typically named with the chemical system (eg "tc-KFMASH") and contain the activity-composition relationships, though they can have any name you choose. Form:

From	What it does
"axfile <name>"</name>	
"axfile ask"	prompts for a name

Default: "axfile ask"

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setdefPwindow

This script sets a default P window to be used if the user hits return (ie inputs nothing) at a prompt for a P (or PT) range. The 2 numbers "defplow" and "defphigh" defining the window must be given. Form:

From	What it does
"setdefPwindow defplow defphigh"	the given numbers define the default P window;
"setdefPwindow yes defplow defphigh"	as above

"setdefPwindow ask defplow defphigh"	you are prompted to determine if the specified window is to be used;
"setdefPwindow no defplow defphigh"	the specified window is ignored

eg "setdefPwindow 2 8"

Default: 0.1 to 20 kbar

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setdefTwindow

This script sets a default T window to be used if the user hits return (ie inputs nothing) at a prompt for a T (or PT) range. The 2 numbers "deftlow" and "defthigh" defining the window must be given. Form:

From	What it does
"setdefTwindow deftlow defthigh"	the given numbers define the default T window;
"setdefTwindow yes deftlow defthigh"	
"setdefTwindow ask deftlow defthigh"	you are prompted to determine if the specified window is to be used;
"setdefTwindow no deftlow defthigh"	the specified window is ignored

eg "setdefPwindow 400 800"

Default: 200 to 1000°C

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setPwindow

This script allows the P window within which equilibria are considered to be specified in the datafile. The 2 numbers "plow" and "phigh" defining the window must be given. With these "phigh" ≥ "plow", with, for example, equality not being allowed if the calculated equilibria are invariant. If equality is specified, and is not allowed, THERMOCALC prompts for a proper range. For calculations at a fixed P (eg for a T-x pseudosection) the script can be followed by only a single number or the same number twice Form:

Form	What it does
"setPwindow plow phigh"	the given numbers define the P window;
"setwindow yes plow phigh"	as above
"setPwindow ask plow phigh"	you are prompted to determine if the specified window is to be used;
"setPwindow no plow phigh"	the specified window is ignored

eg "setPwindow 2 8" to calculate in this range or "setPwindow 8 8" to calculate at fixed P

Default: prompt for P window.

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setTwindow

This script allows the T window within which equilibria are considered to be specified in the datafile. The 2 numbers "tlow" and "thigh" defining the window must be given. With these "thigh" ≥ "tlow", with, for example, equality not being allowed if the calculated equilibria are invariant. If equality is specified, and is not allowed, THERMOCALC prompts for a proper range. For calculations at a fixed T (eg for a P-x pseudosection) the script can be followed by only a single number or the same number twice Form:

Form	What it does
"setTwindow tlow thigh"	the given numbers define the T window;
"setwindow yes tlow thigh"	as above
"setTwindow ask tlow thigh"	you are prompted to determine if the specified window is to be used;
"setTwindow no tlow thigh"	the specified window is ignored

eg "setTwindow 200 800" to calculate in this range or "setTwindow 600 600" to calculate at fixed T

Default: prompt for T window.

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calctatp

With a PT window (see setwindow), and considering univariant equilibria (ie reaction lines) in a PT window, there is a choice of whether T should be calculated at a series of P, or P calculated at a series of T. For example, for reactions with a very small dP/dT, it is easier to calculate P at T. This script allows this choice to be made. (If for the window plow=phigh or tlow=thigh, then the choice does not arise). Form:

Form	What it does
"calctatp"	always calculate T at P
"calctatp yes"	as above
"calctatp ask"	you are prompted for the choice
"calctatp no"	always calculate P at T

Default: you are prompted for the choice

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moreprec (defunct in version 3.30)

Allows an extra decimal place in the output of PT and x values. This is useful if equilibria are closely spaced - note, however, that there is no implication that the numbers are somehow more accurate when this switch is on. Form:

Form	What it does
"moreprec"	an extra decimal place is used in the output;
"moreprec yes"	as above
"moreprec ask"	you are prompted whether an extra decimal place should be used in the output;
"moreprec no"	no extra decimal place in the output.

Default: "moreprec no"

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calcsdnle

Allows uncertainty propagation calculations to be switched on or off. Note the uncertainties given are 1sigma, but should be quoted at 2sigma (95% confidence). Form:

Form	What it does
"calcsdnle"	uncertainty propagation calculations will be performed
"calcsdnle yes"	
"calcsdnle ask"	you are prompted whether uncertainty propagation calculations will be performed;
"calcsdnle no"	no uncertainty propagation

Default: "calcsdnle no"

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omit or ignore

Allows phases for which information is provided in the first part of a datafile to be ignored in using the datafile. This circumvents the need for lengthy lists of phases to be omitted at the 'which phases' prompt when running THERMOCALC or alternatively the cutting and pasting of phase info from the first part to the last (storage) part of datafiles. Form:

Form	What it does
"ignore phase1 phase2 etc"	phase1, phase2, etc, which are present in the first part of the datafile, are to be ignored in the calculations
"ignore yes phase1 phase2 etc"	as above
"ignore ask phase1 phase2 etc"	you are prompted whether these phases are to be ignored
"ignore no anything"	no phases ignored
"ignore ———"	nothing following the script means no phases ignored

Default: nothing ignored

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fluidpresent

In phase diagram calculations involving H2O or CO2 (but not H2O and CO2, see below), these end-members might be included in a system to be used in fluid-absent equilibria (say with a(H2O) set at particular values). This script is used to communicate that a system may have H2O or CO2 present as phases. If a fluid phase with H2O and CO2 is in the system, it is coded in the datafile as a solution, much as solid solutions are, and mode 1 should be used. In this case fluidpresent is automatically yes, and does not need to be scripted. Form:

Form	What it does
"fluidpresent"	H2O or CO2 is present as a phase;
"fluidpresent yes"	as above
"fluidpresent ask"	prompt - this is THERMOCALC normal behaviour in the absence of the script;
"fluidpresent no"	H2O or CO2 cannot be present as a phase

Default: "fluidpresent ask"

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fluidexcess

In calculations, it is often convenient to consider one or more phases to be "in excess", to reduce the total number of equilibria to be considered, and also to reduce the effective dimension of the system so that compatibility diagrams can be drawn (eg the 6-component system KFMASH is reduced to the effective ternary, AFM, by having muscovite, quartz and H2O in excess). The fluidexcess script allows fluid in excess to be specified. Form:

Form	What it does
"fluidexcess"	fluid is in excess;
"fluidexcess yes"	as above
"fluidexcess ask"	prompt - this is THERMOCALC normal behaviour in the absence of the script;
"fluidexcess no"	fluid not in excess

Default: "fluidexcess ask"

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setexcess

In calculations, it is often convenient to consider one or more phases to be "in excess", to reduce the total number of equilibria to be considered, and also to reduce the effective dimension of the system so that compatibility diagrams can be drawn (eg the 6-component system KFMASH is reduced to the effective ternary, AFM, by having muscovite, quartz and H2O in excess). The setexcess script allows phases other than fluid to be considered to be in excess (see fluidexcess). Form:

Form	What it does
"setexcess ph1 ph2 ph3 "	the phases ph1, ph2, and so on are taken to be in excess (above eg "setexcess mu q");
"setexcess yes ph1 ph2 ph3"	as above
"setexcess ask ph1 ph2 ph3"	prompt to determine if the phases ph1, ph2, and so on are taken to be in excess
"setexcess no ph1 ph2 ph3"	nothing in excess

Default: prompt for in excess phases.

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exbuff

Allows external buffering calculations involving H2O-CO2 fluids to be done. For example in calculating a T-x(CO2), incombination with "setiso xCO2", it allows pseudosections to be drawn. Form:

Form	What it does
"exbuff"	external buffering is activated
"exbuff yes"	as above
"exbuff no"	not external buffering.

Default: "exbuff no"

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seta

In phase diagram calculations, it is sometimes important to see what the consequence is of varying the activity of an end-member, for example H2O (under granulite facies conditions). This script allows this to be done. Form:

Form	What it does
"seta"	you will be prompted for the name of the end-member whose activity will be fixed;
"seta yes"	as above
"seta em"	the activity of end-member, em, will be fixed (eg "setaH2O");
"seta yes em"	as above
"seta ask"	you will be prompted to determine if you wish to fix an activity;
"seta ask em"	you will be prompted to determine if you wish to fix the activity of end-member, em (eg "seta ask H2O");
"seta no anything"	you do not wish to fix an activity.

Default: "seta no"

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setmu

In phase diagram calculations, it is sometimes important to see what the consequence is of varying the chemical potential of a component, for example K2O (in orthoamphibole assemblages). This script allows this to be done. This script is also used for construction of μ - μ diagrams via setting a range of μ 's to calculate lines Form:

Form	What it does
	you will be prompted for the name of a component whose chemical potential will be fixed

"setmu"	and later, the chemical potential value or range of values and an increment
"setmu yes"	as above
"setmu comp"	the chemical potential of component, comp, will be fixed (eg "setmu Na2O");
"setmu yes comp"	as above
"setmu comp1 num1"	the chemical potential of component, comp, will be fixed at the defined value (eg "setmu Na2O -960"). More than one component may have a fixed mu (eg "setmu yes K2O -939.6960 Al2O3 -1751.0310")
"setmu yes comp1 num1"	as above
"setmu ask"	you will be prompted to determine if you wish to fix a chemical potential;
"setmu ask comp"	you will be prompted to determine if you wish to fix the chemical potential of comp (eg "setmu ask Na2O")
"setmu no anything"	you do not wish to fix a chemical potential.

Default: "setmu no"

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usecalcq

This script prompts THERMOCALC to solve the state of order in order-disorder minerals prior to solving for the specified equilibria. The state of order calculated is then used as a starting guess. The state of order is solve for the P-T condition in the middle of the P-T window specified. Form:

Form	What it does
"usecalcq"	the state of order pre- calculated for order-disorder phases

"usecalcq yes"	as above
"usecalcq ask"	you will be prompted whether you wish the state of order to be pre-calculated
"usecalcq no"	the state of order pre- calculated

Default: "usecalcq yes"

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samecoding

This script causes output to be lined up for visual comparison. All the samecoding phases need to be at the beginning of the axfile. (This is a precursor to just putting the a-x relations in once each for amph, cpx etc). This is currently being debugged - if you are having problems with it, just don't use it (and maybe report to RP what happens). Form:

Form	What it does
"samecoding min1 min2 min3"	causes output to be lined up for visual comparison.

eg to use same coding for coexisting amphiboles and coexisting cpx

samecoding act hb gl samecoding jd o di

on separate lines like this.

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readxyz (version 3.30 and above)

This script followed by a composition variable and a numeric value allows the starting guesses to be set in the scripts file. Doing so makes these starting guesses overwrite those in the a-x file for each calculation. A THERMOCALC-readable list can be printed in the tc-log file using the "printxyz" script (see above). This is an easy way to change the staring guesses without having to open the a-x file Form:

Form	What it does
"readxyz var num"	this sets this number as the starting guess for the chosen variable (eg "readxyz x(bi) 0.4284")

Default: "script not present"

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pseudosection

In phase diagram calculations, it is very informative to know if a particular equilibrium is "seen" by a particular bulk composition. Phase diagrams, for example PT diagrams, drawn to show just those equilibria "seen" by a particular bulk composition are referred to as pseudosections. This script allows pseudosection information to be collected. Typically the bulk composition is set using the "setbulk" or "rbi" script but can be input manually at each calculation. Form:

Form	What it does
"pseudosection"	you are prompted for the bulk composition(s) you want to collect pseudosection information for
"pseudosection yes"	as above
"pseudosection ask"	you are prompted concerning whether you want to collect pseudosectio information; if you reply yes, you are prompted for the bulk composition(s) you want to collect pseudosection information for; if you reply no, pseudosection informationis not collected;
"pseudosection no"	pseudosection information is not collected.

Default: "pseudosection ask"

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setbulk

If pseudosection is yes, then setbulk allows one or two bulk compositions (in mole %) to be scripted. The setbulk script is followed by the bulk composition in the system defined by the a-x models. Each bulk composition has the components in a specific order. The

simplest way to get this order correct is to run THERMOCALC with the datafile of interest, with "pseudosection yes" and without the setbulk script. The bulk composition is then prompted for in the correct order. The bulk compositions input do not have to sum to 100, as this normalization is done by THERMOCALC. Each component in the system (including those only in inexcess phases) must be defined with the exception of fluid. Forms:

Form	What it does
"setbulk yes com1 com2 com3"	use this bulk composition (com = component)
"setbulk com1 com2 com3"	
"setbulk com1 com2 com3"	prompt to ask the user whether this bulk composition is to be used
"setbulk no com1 com2 com3"	this bulk composition is not to be used, and the user will be prompted for the bulk composition (useful for temporarily "switching off" the script)

An example of how this script looks in a typical P-T pseudosection file is below:

An example of how this script looks in a typical P-x ot T-x pseudosection file is below:

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setmodeiso

In pseudosection calculations, a useful facility is to be able to fix the value of modes of phases, and calculate all the other composition and mode variables of the phases in an equilibrium. This facility allows the position of mode isopleths on a PT pseudosection to be calculated. In particular, it allows (in conjunction with the "zeromodeiso" script) the boundaries between fiels on a pseudosection to be calculated. Regardless of the variance of the field in a PT pseudosection, with one mode fixed, the equilibrium is effectively univariant (ie defines a PT line), and with two modes fixed, the equilibrium is effectively invariant (ie defines a PT point). Form:

Form	What it does
"setmodeiso"	you will be prompted for the names of the composition variables to be fixed;
"setmodeiso yes"	as above
"setmodeiso ph"	the mode of the phase, ph, is to be fixed, and the values are prompted for (eg "setmodeiso bi");
"setmodeiso yes ph"	as above
"setmodeiso ph isolow isohigh isoinc"	the mode of the phase, ph, is to have values running from isolow through to isohigh with an interval, isoinc (eg "setiso bi 0.1 0.6 0.1")
"setmodeiso yes ph isolow isohigh isoinc"	as above
"setmodeiso ask"	you will be prompted to determine if you wish to fix any composition variables, then prompted for the details;
"setmodeiso ask ph"	you will be prompted to determine if you wish to fix the mode of the phase, ph (eg "setmodeiso ask bi");
"setmodeiso ask ph isolow isohigh isoinc"	you will be prompted to determine if you wish to fix the composition variable, ph, and to have values running from isolow through to isohigh with an interval, isoinc;
"setmodeiso no anything"	you do not wish to fix any modes

Default: "setmodeiso ask"

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zeromodeiso

This qualifies "setmodeiso yes", making the setting of modes always involve zero values. This facility is central to calculating the lines and points that define filed boundaries in PT and Tx pseudosections. Form:

Form	What it does
"zeromodeiso"	allows zero mode isopleths to be set
"zeromodeiso yes"	as above
"zeromodeiso ask"	you are prompted whether you want zero mode isopleths to be set
"zeromodeiso no"	zero mode isopleths are not set

Default: "zeromodeiso ask"

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setiso

In phase diagram calculations, an important facility is to be able to fix the value of any compositional variable, and calculate all the other composition variables of the phases in an equilibrium. This facility allows the position of isopleths on a PT diagram to be calculated, as well as the edges of trivariant fields on compatibility diagrams. For example, in AFM (in KFMASH), the compositions of, say, coexisting biotite and chlorite can be calculated via fixing x(Fe) in the biotite (or chlorite). A major application concerns Tx(CO2) (and P-x(CO2)) diagrams: by fixing x(CO2) in the fluid and calculating equilibria such diagrams can be calculated. Care is needed in the specification of variance for such calculations. Remember that univariant lines on Tx(CO2) diagrams are actually divariant on a PT diagram. Thus with x(CO2) to be specified, the required variance when prompted to calculate univariant lines on a Tx(CO2) is trivariant. In general, for each composition variable fixed, the (effective) variance is decreased by one. Form:

Form	What it does
	you will be prompted for the

"setiso"	names of the composition variables to be fixed;
"setiso yes"	as above
"setiso var"	the composition variable, var, is to be fixed, and the values are prompted for (eg "setiso x(bi)"); "setiso var isolow isohigh isoinc"
"setiso yes var"	as above
"setiso yes var isolow isohigh isoinc"	the composition variable, var, is to have values running from isolow through to isohigh with an interval, isoinc (eg "setiso x(bi) 0.1 0.6 0.1")
"setiso ask"	you will be prompted to determine if you wish to fix any composition variables
"setiso ask var"	you will be prompted to determine if you wish to fix the compositional variable, var (eg "setiso ask x(bi)")
"setiso ask var isolow isohigh isoinc"	you will be prompted to determine if you wish to fix the composition variable, var, to have values running from isolow through to isohigh with an interval, isoinc
"setiso no anything"	you do not wish to fix any composition variables

Default: "setiso ask"

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calcg

The Gibbs energy of each equilibrium is calculated. Form:

Form	What it does
"calcg"	calculate Gibbs energy
"calcg yes"	as above
"calcg ask"	prompt for calculation of Gibbs energy

"calcg do not calculate Gibbs energy no"

Default: "calcg no"

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dogmin

Allows Gibbs energy minimisation calculations: for a specified bulk composition all of the equilibria from divariant up to a specified variance are calculated and the one, at each PT, that has the lowest Gibbs energy is the stable one. This can be useful for kick-starting the calculation of a pseudosection, but in large chemical systems can be very slow. Form:

Form	What it does
"dogmin"	do Gibbs energy minimisation
"dogmin yes"	as above
"dogmin ask"	prompt for Gibbs energy minimisation
"dogmin no"	Gibbs energy minimisation not done

Default: "dogmin no"

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rbi

This script allows the bulk rock composition to be defined via a combination of mineral compositions and modes. This can be useful for adding/removing a proportion of a given phase (eg to model melt loss). A readable table of mineral compositions and modes can be printed into the tc-log file using the script "printbulkinfo" Form:

the table of mineral compositions will look something like this:

```
%
%
                   cd
liq ksp q
rbi yes 0.128725 0.142898
0.590856
                         0.137521
%
                  H20
                              Si<sub>02</sub>
A1203
               Ca0
                            Mg0
             K20
                        Na<sub>20</sub>
Fe0
           0.355525 5.000000
                                      2.000000
rbi
```

```
1.245074
                   0.754926
              % cd
rbi
               0 3.000000
                            1.000000
0.021669 0.750962 2.227370
              % g
         0
        0.320534 2.252573 0.216288
rbi
0.012426 0.020572 0.070052 0.149223
           % liq
0.029450
                  2.993191 0.503404
rbi
0.006809
                0
                         0 0.458251
0.038344
           % ksp
                  1.000000
rbi
0
         0
                             0
              % q
0
         0
%
```

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calcah2o

This script allows the activity of water to be calculated for each calculation. This is useful to investigate variations in aH2O around a diagram Form:

Form	What it does
"calcah2o"	the activity of water will be calculated
"calcah2o yes"	as above
"calcah2o ask"	you will be prompted whether you want the activity of water to be calculated
"calcah2o no"	the activity of water will not be calculated

Default: "calcah2o no"

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calcmu

This script allows the chemical potential of each component to be calculated for each calculation. This is used for construction of μ - μ diagrams. The chemical potential values will also be printed in the tc-<filename>-dr.txt file. Form:

Form	What it does	
"calcmu"	the chemical potentials will be calculated	
"calcmu yes"	as above	
"calcmu	you will be prompted whether	

ask"	you want the chemical potentials to be calculated
"calcmu no"	the chemical potentials will not be calculated

Default: "calcmu no"

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drawpd

If "makeo yes" is in the preferences file, indicating that a 'other' file is to be made (whose default name is 'tc-<filename>-dr.txt'), this causes coordinate information from the calculations to be put in it in a format that is readable by the drawpd software. Form:

Form	What it does
"drawpd"	put coordinate info in 'other' file
"drawpd yes"	as above
"drawpd no"	nothing put

Default: "drawpd no"

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project

In representing the results of phase diagram calculations, it is often convenient to be able to project the calculated compositions of the phases into a compatibility diagram. This script allows this. Form:

Form	What it does
"project"	you are prompted for the definition of the compatibility diagram you want to use
"project yes"	as above
"project ask"	you are prompted concerning whether you want to project the phase diagram information; if you reply yes, you are prompted for the definition of the compatibility diagram you want to use; if you reply no then projection is not performed

"project projection is not performed.
no"

Default: "project ask"

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setproject

If project is yes, then setproject allows the definition of the compatibility diagram (or projection plane) to be scripted, at least for compatibility diagrams which have oxides as their corners. So for example, for KFMASH projected from muscovite, quartz and H2O (ie having muscovite, quartz and H2O in excess), Al2O3, FeO and MgO (ie AFM) define the corners of the usually used compatibility diagram. The oxides are specified in terms of their position in the list of oxides given by THERMOCALC when this same information

is prompted for. The simplest way to get this script correct is to run THERMOCALC with the datafile of interest, with "project yes" and without the setproject script. Forms:

Form	What it does
"setproject yes 1 3 2"	use 1st, 3rd and 2nd component
"setproject 1 3 2"	as above
"setproject ask 1 3 2"	prompt to ask the user whether this is to be used
"setproject no 1 3 2"	this is not to be used, and the user will be prompted for the projection plane (useful for temporarily "switching off" the script)

below is the setproject script for a standard AFM diagram

%			
setproject	٨	1	۵
0 0	^	_	O
setproject	F	0	0
1 0	М	0	1
setproject yes 0 0	[V]	Ø	T
%			

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projcomp

In phase diagram calculations involving projection and the collection of pseudosection information, particularly when there is projection from variable composition phases, it can be useful to see where the bulk composition(s) plot on the compatibility diagram (projection plane). This script allows this, as long as project is yes and pseudo is yes. A bulk composition is only projected if the calculated equilibrium is "seen" by the bulk composition. Form:

Form	What it does
"projcomp"	do project bulk composition(s) onto the compatibility diagram
"projcomp yes"	as above
"projcomp ask"	you are prompted whether you want bulk composition(s) projected onto the compatibility diagram
"projcomp no"	bulk composition(s) are not projected

Default: "projcomp ask"

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printbulkinfo

This script causes the bulk rock composition to be printed in the tc-log file as a THERMOCALC-readable table of mineral modes and mineral compositions such as the table above for each P-T calculation. This can then be copied and pasted into the script file. Form:

Form	What it does
"printbulkinfo"	the bulk rock composition table is printed in the tc-log file
"printbulkinfo yes"	as above
"printbulkinfo ask"	you will be prompted whether you want this table printed
"printbulkinfo no"	the bulk rock composition table is not printed in the tc-log file

Default: "printbulkinfo no"

printguessform (version 3.23-3.26).

This script causes the mineral composition results from each calculated point to be printed in the tc-log file as a list of mineral compositions in the form of the composition variables for each phase. This can then be copied and pasted into the datafile as the starting guesses. Form:

Form	What it does
"printguessform"	the mineral composition list is printed in the tc-log file
"printguessform yes"	as above
"printguessform ask"	you will be prompted whether you want this list printed
"printguessform no"	the mineral composition list is not printed in the tc-log file

The list will look something like this:

x(cd)	0.3775
h(cd)	0.3555
x(g)	0.7425
m(g)	0.2503
$q(\tilde{L})$	0.2713
fsp(L)	0.3573
na(L)	0.1648
an(L)	0.01243
ol(L)	0.02266
x(L)	0.7730
h2o(L)	0.3205
na(ksp)	0.07669
ca(ksp)	0.006809

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printxyz (version 3.30 and above)

This script causes the mineral composition results from each calculated point to be printed in the tc-log file as a THERMOCALC-readable list of mineral compositions in the form of the composition variables for each phase. This can then be copied and pasted into the script file as the starting guesses. Doing so makes these starting guesses overwrite those in the a-x file. Form:

Form	What it does
"printxyz"	the mineral composition list is printed in the tc-log file

"printxyz yes"	as above
"printxyz ask"	you will be prompted whether you want this list printed
"printxyz no"	the mineral composition list is not printed in the tc-log file

Default: "printxyz no"

The list will look something like this:

```
% calculated at P =
                           6.9; T =
                                       800
readxyz q(L)
readxyz fsp(L)
                           0.1446
                           0.3061
readxyz na(L) readxyz an(L)
                           0.6839
                          0.01097
readxyz ol(L)
                        0.009983
readxyz x(L)
                           0.7444
readxyz h2o(L)
                           0.5197
readxyz x(bi)
                          0.4284
readxyz y(bi)
readxyz f(bi)
                          0.2536
                          0.03740
readxýz t(bi)
                          0.1263
readxyz Q(bi)
                          0.2588
readxyz x(g)
                          0.7238
readxyz z(g)
readxyz f(g)
readxyz ca(pl)
readxyz k(pl)
                         0.02970
                        0.005363
                         0.3011
                          0.03034
```

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incthermo

In phase diagram calculations, there is the option of a table of the thermodynamic data being included in the logfile. This script allows it to happen. Usually there is not much point in including the table, given that it takes a little time to print. One situation it might be useful, though, is when the DQF facility or the dependent endmember facility is used, in which case the data for the constructed end-member may be of interest. Form:

Form	What it does
"incthermo"	include the table
"incthermo yes"	as above
"incthermo no"	table not included

Default: "incthermo no"

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incax

In phase diagram calculations, there is the option of a summary of the activity-composition relationships in the datafile being included in the logfile. This script controls this. Given that coding the activity-composition relationships provides lots of opportunities for making mistakes, it is a good idea to include this information at least until it has been thoroughly checked. Form:

Form	What it does
"incax"	include the a-x info;
"incax yes"	as above
"incax no"	a-x info not included

Default: "incax yes"

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incnler

In phase diagram calculations, there is the option of the set of reactions between the end-members of the minerals used in setting up the set of non-linear equations (equilibrium relationships) being included in the logfile. This script allows it to happen. Usually there is not much point in including this information. Form:

Form	What it does
"incnler"	include the reactions;
"incnler yes"	as above
"incnler no"	reactions not included

Default: reactions not included

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Isqds

This script allows activities of end-members rather than mineral comps to be output for calculations. Form:

Form	What it does
"Isqds"	end member activities output
"Isqds yes"	as above
"Isqds no"	mineral compositions output

Default: "Isqds no"

The output will look like this:

```
P(kbar) T(?C) kspL

anL silL foL faL

h2oL phl

8.80 866.2 0.169 0.00327

0.0398 1.18e-5 0.00215 0.250

0.0659

mode liq bi g

ksp ilm sill q

0.3607 0 0.2377

0.09587 0.02994 0.02508 0.2508
```

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smath

If "makeo yes" is in the preferences file, indicating that a dr file is to be made (whose default name is tc-<name>-dr), this causes coordinate information from the calculations to be put in it. The default form of the coordinate information is Mathematica-readable (ie info being comma-separated, structured with curly brackets). This is the standard information used with the Mathematica "drawline" code for generating various diagrams. This can be varied, or changed via, "xinsmath yes" or "tabsmath yes". Form:

Form	What it does
"smath"	put coordinate info in dr file;
"smath yes"	as above
"smath no"	nothing put

Default: "smath no"

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xinsmath

This qualifies "smath yes". Generally the coordinates output are just the PT. For some calculations, eg for T-x(CO2) diagrams, composition info is also needed. This script causes it to be output in addition to the PT info. Form:

Form	What it does
"xinsmath"	put extra coordinate info in "other" file
"xinsmath yes"	as above

"xinsmath no" nothing put

Default: "xinsmath no"

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tabsmath

This qualifies "smath yes". Rather than the coordinate information being in Mathematica-readable format, it is output in simple tab-separated form (as understood by various spreadsheet programs). Form:

Form	What it does
"tabsmath"	do tab-separated output;
"tabsmath yes"	as above
"tabsmath ask"	ask whether to produce tab- separated output;
"tabsmath no"	output not tab-separated

Default: "tabsmath no"

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cmath

If "makeo yes" is in the preferences file, indicating that a 'other' file is to be made (whose default name root is 'th dr' - Win: 'thdr.txt'), this causes compatibility diagram coordinate information from the calculations to be put in it. The form of the coordinate information is Mathematica-readable (ie info being comma-separated, structured with curly brackets), and is intended for use with the Mathematica functions for drawing compatibility diagrams. Form:

Form	What it does
"cmath"	put coordinate info in "other" file
"cmath yes"	as above
"cmath no"	nothing put

Default: "cmath no"

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txmath

If "makeo yes" is in the preferences file, indicating that a 'other' file is to be made (whose default name root tc-<name>-dr), this causes Tx/Px pseudosection coordinate information from the calculations to be put in it. The form of the coordinate information is Mathematica-readable (ie info being comma-separated, structured with curly brackets), and is intended for use with the Mathematica functions for drawing these pseudosections. Form:

Form	What it does
"txmath"	put coordinate info in "other" file
"txmath yes"	as above
"txmath no"	nothing put

Default: "txmath no"

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scripts for avPT (mode 2)

defsdaparams

for setting activity uncertainty defaults

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usedefsda

using the defaults

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justrelu

just relative uncertainties (as in the thermobarometry paper)

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setrockname

allows the results to be labelled with the rock name

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diagnostics

to always see the diagnostics

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scripts for calculations not involving (solid) solutions (mode 3)

setmu

allows you to set the chemical potentials of components

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seta / setlna / setloga

allows you to set activities of components

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