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Geochemical Data Toolkit (*GCDkit*) written in R language

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<http://www.gcdkit.org>

**User's Guide to version 6.2
(12 January 2024)**



GCDkit

is a system for handling and recalculation of whole-rock analyses from igneous rocks. It is fully menu driven but, at the same time, can be used in an interactive regime (power users only).

Using *GCDkit*, the data can be loaded, individual analyses grouped into coherent groups or searched according to various criteria. They can be plotted into commonly used classification diagrams (e.g. Harker plots, TAS, AFM, R_1 – R_2) as well as a variety of user-defined plots (binary and ternary plots, multiple plots, spider plots).

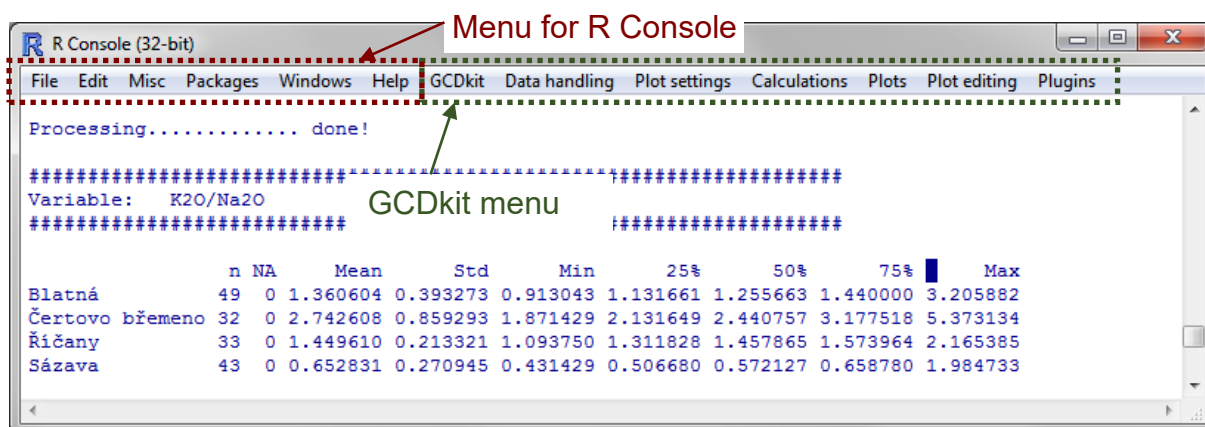
Moreover, some basic statistical methods are implemented (both for the whole data set and individual groups) using a fraction of the power built in R language. Included are the descriptive statistics, boxplots, histograms, principal component and cluster analysis. *GCDkit* also makes possible the most common normative calculations to be performed.

The system is easy to expand by means of the user-defined diagrams or so-called plugins that provide a simple method of adding effortlessly new items to the menus of *GCDkit*. Nine of the latter are included in the current distribution, namely modules for interpretation of the radiogenic isotope data (*SrNd.r*: calculating Sr and Nd initial compositions, Nd model ages, plotting isochrons etc.), modules estimating saturation temperatures of accessory phases (*saturation.r*: zircon, apatite, monazite and rutile), temperatures of granitic magmas production (*JungAlTitemp.r*), isocon/wedge plots (*isocon.r*), a module assessing the magnitude of the REE tetrad effect (*tetrad.r*), *clr* transformation of (closed) compositional data (*disclosure.r*), *escoRt* – the implementation of the ESCORT expert system (Pearce 1987) to identify geodynamic setting of igneous rocks and the “projection from biotite” of Moyen *et al.* (2017). The last one is an online interface to the EarthChem database (see below).

Important!

This *GCDkit* was built in **R version 4.1.3** for Windows. Also this manual applies in its entirety only to the Windows (GUI) version of *GCDkit*. Due to the quick developments in the R project, it is blocked in any other versions!

The platform-independent (i.e. non-Windows) mutation of *GCDkit* is still experimental. Some of the features may not be available, others may work unpredictably. Please use sensibly, at your own risk. For further information, see <http://blog.gcdkit.org>.



TECHNICAL NOTES

- After starting R opens “R Console”, a text window serving for entry of commands as well as display of textual output. In addition, during the session are typically opened one or more graphical windows.
- Please note that on the Windows systems, the *GCDkit* menu appears right of the menus of the R system itself (see Figure above). On non-Windows systems, please type *menuet()* into the R command prompt in order to start the Tcl/Tk menu system.
- The interactions built in many plots (most commonly data point identification) can be stopped from the menu that appears after pressing the right mouse button.
- If experiencing problems, the computation/plotting can be interrupted anytime by hitting *Escape* key or from the menu (*Misc|Stop current computation*).
- The errors encountered while running the *GCDkit* system and displayed in the Console are (mostly) not fatal. In most cases it means that the command can be re-run (one can simply press the up/down arrows in the Console to scroll through the commands history), entering/modifying the parameters that caused the crash. If the problem persists, record the details (including the version of *GCDkit*, R and your operation system) and send the data file to us, so we can track down and fix the bug.
- *GCDkit* requires the R packages *compiler*, *foreign*, *graphics*, *grDevices*, *grid*, *IRdisplay*, *lattice*, *MASS*, *stats* and *utils* to be installed (but this is normally the case).
- Moreover, the package *sp* has to be always present as it is used for classification purposes, and the package *R2HTML* that provides output to HTML format. The package *XML* allows access needed to read input from online databases. On Windows, highly recommended is also the package *RODBC* enabling import and export into the MS Excel, Access and dBase (DBF) formats. Packages *curl*, *rgdal* and *tkrplot* are utilized solely for the online access to the EartChem database. All these packages can be installed from our distribution. But if need be, any of the above packages can be added via the menu *Packages|Install package(s)...* (you need to be online). Offline installation is also possible, though: zip file downloaded from the CRAN site (<http://www.r-project.org>) can be introduced via menu *Packages|Install package(s) from local zip files...*
- **Note that recommended systems** to run *GCDkit* are Windows 11/10/8/7. The current version should also work on Mac OS X (release 10.6 and above) and various distributions of Linux (Debian, RedHat, SUSE, Ubuntu).
- It is always a good idea to close the unnecessary graphical windows, for instance using the function *graphicsOff()* or the corresponding item from the menu *GCDkit*.
- By the same token, please try to remove periodically all unwanted datasets still stored in the memory. For this purpose serves the function *purgeDatasets()* or the corresponding item from the *GCDkit* menu.

- More immediate response is obtained when print buffering (*Misc|Buffered Output*) is disabled.
- For graphical output, *PDF* format should be given preference, as it is the most accurate and it also supports semitransparency used in some diagrams. If no semi-transparent colours are desired, PostScript is a viable alternative. Using *WMF* (Windows metafiles) or copying via clipboard are to be discouraged, as the latter two methods lead sometimes to distortions of the graphs. For saving, simply right-click the graphical window and select the desired format.
- If semi-transparent colours are used, some printers simply ignore them when printing directly from the plotting window. In these cases, use PDF export first and then print from some other application, such as Adobe Reader.
- The use of special symbols or accented characters (such as in some East European languages) in the datasets is mostly okay but it may also sometimes cause unexpected problems. So please be sensible, especially in the variable names.

Quick guide to data files

GCDkit requires plain text data files, delimited by tabs, commas or semicolons (delimiter is recognised automatically, as is a decimal point/comma). See the testing files *sazava.data* and *cbp_iso.data* that come with the *GCDkit* distribution for examples of valid data files with whole-rock geochemical and Sr–Nd isotopic data, respectively.

As an (less forgiving) alternative on operation systems with graphical user interface, the data can be pasted, via clipboard, from any well-behaved software, including Excel. Be careful, though, as the precision of the numbers imported will be that *displayed* in the original Excel file!

If library *RODBC* has been installed, and R is run in **32-bit mode**, the *GCDkit.Mineral* first attempts to establish ODBC connection to the selected file, and open it as a Excel (*.XLS, *.XLSX) or Access (*.MDB) format.

Without ODBC support, the *GCDkit* can read plain text files, including the variants represented by the *NewPet* (*.ROC) (Clarke *et al.*, 1994) or *PetroGraph* (*.PEG) (Petrelli *et al.*, 2005) and the outputs from World Wide Web-based databases such as *GEOROC* (<http://georoc.mpch-mainz.gwdg.de/georoc>), *EarthChem* and *PETDB*, both hosted at the same web address (<http://www.earthchem.org>). Available is also import of comma-separated text (*.CSV) and dBase III/IV (*.DBF) files. The latter were invoked to store data by the *MinPet* package (Richard, 1995), once used mainly for mineral recalculations. It, however, runs only on now obsolete 16-bit Windows. Moreover, the DBF files are/have been used to store data by other popular geochemical packages, such as *IgPet* (Carr & Gazel, 2017), and the output of the online *NAVDAT* database (<http://www.navdat.org>).

On Windows, there a possibility of searching the *EarthChem* online database (<http://www.earthchem.org>) directly from within the *GCDkit* menu.

As a rule of thumb, if encountering any problems with loading data, please use import from text file or *via* clipboard, the methods that are much more robust than any other one.

In text files and spreadsheets, the first row should contain names of the data columns (except for the first column that is automatically assumed to contain sample names). Hence the first line of the text file may (or may not) have one item less than the following ones.

Missing values (NA) are allowed anywhere in the data file body; values equalling to zero, or any of 'NA', 'N.A.', '-', 'b.d.', 'bd', 'b.d.l.', 'bdl' and 'n.d.' are also treated as such. While

loading Excel files, the values *#WHATEVER!* (Excel error messages) are also replaced by ‘NA’.

If desired, negative numbers and values ‘< x’ (used by some authors to indicate items below detection limit) can be either replaced by their half (i.e. interpreted as a half of the detection limit) or ‘NA’. User is prompted which of these options he prefers.

Alternatively, negative values can be left untouched (as may be desirable e.g. for $\delta^{18}\text{O}$ values).

The data rows start with sample name and do not have to be all of the same length (the rest of the row is filled by ‘NA’ automatically).

A column named “**Symbol**” (if any) is taken as containing specification of plotting symbols. They can be specified either as single character strings or numeric codes, whose overview is given on the right or can be obtained by invoking the menu item *Data handling|Show available symbols*.

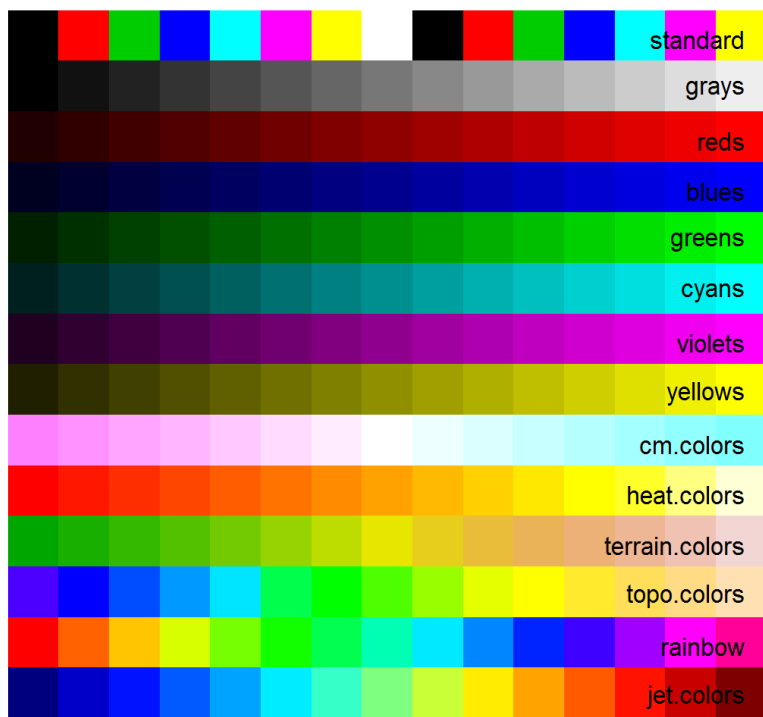
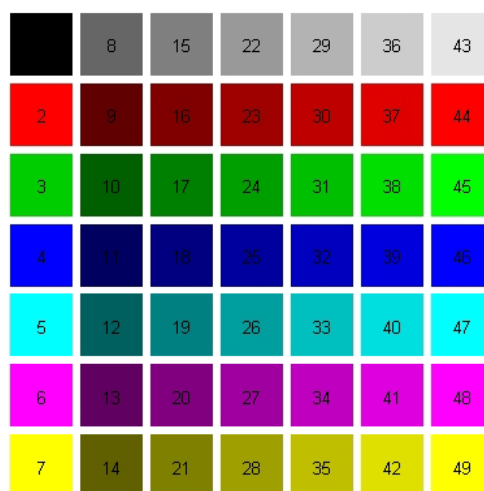
Another column, whose name is “**Colour**” or “**Color**” (if any, capitalization does not matter) may contain codes (1–49) or English names for the plotting colours. There are 657 of the latter, see Help, invoke menu *Data handling|Show available colours* or type *colours()* into the *R Console*.

Alternatively, colours can be coded using the RGB model (8-bit hexadecimal colour code) (e.g. *#FF0000* for red) or RGBA model (the same but with two extra hexadecimal digits appended for the alpha channel, expressing the colour’s opacity). For instance, *#FF0000BF* is red with 75% opacity (~ 25% transparency).

If specifications of both the plotting symbols and colours are missing completely, and at least one non-numeric variable is present, the user is prompted whether he does want to have the symbols and colours assigned automatically, according to individual values (levels) of the selected label. Otherwise default symbols (empty black circles) are used.

A column named “**Size**” or “**cex**” may contain relative size for each of the plotting symbols (for instance, use 1 for a standard size, 2 for double and 0.5 for half the size).

Apostrophes and any other special characters should be avoided in the sample/variable names. Else, the data files are



practically freeform, i.e. no specified oxides/elements are required and no exact order of these is to be adhered to. Analyses can contain as many numeric columns as necessary; the names of oxides and trace elements are self-explanatory (e.g. 'SiO2', 'Fe2O3', 'Rb', 'Nd'). Total iron, if given, should be expressed as ferrous oxide ('FeOt', 'FeOT', 'FeOtot', 'FeOTOT' or 'FeO*') or ferric oxide ('Fe2O3t', 'Fe2O3T', 'Fe2O3tot', 'Fe2O3TOT' or 'Fe2O3*'). Structurally bound water can be named 'H2O.PLUS', 'H2O.P', 'H2O+', 'H2OPLUS' or 'H2O_PLUS'. For isotopic ratios, please use simple forms such as '87Rb/86Sr', '87Sr/86Sr', '147Sm/144Nd', '143Nd/144Nd' or 'delta18O'.

Note that names of variables are **case sensitive** in R. However, any of the fully upper case names of the oxides/elements that appear in the following list are translated automatically to the appropriate capitalization:

SiO2, TiO2, Al2O3, Fe2O3, FeO, MnO, MgO, CaO, Na2O, FeOt, Fe2O3t, Li2O, mg#, Li, Rb, Cs, Be, Sr, Ba, Sc, Y, Ti, Zr, Hf, V, Nb, Ta, Cr, Mo, W, Re, Ru, Os, Co, Rh, Ir, Ni, Pd, Pt, Cu, Ag, Au, Zn, Cd, B, Ga, In, Tl, C, Ge, Sn, Pb, P, As, Sb, Bi, S, Se, Te, F, Cl, Br, I, At, La, Ce, Pr, Nd, Pm, Sm, Eu, Gd, Tb, Dy, Ho, Er, Tm, Yb, Lu.

Also trailing spaces in the end of the column names are disposed of. However, note that, with some exceptions such as importing from online databases, **no duplicated column/sample names are allowed!** The correct syntax is checked upon loading.

After loading, all the completely empty columns are removed first. In the text files, any line starting with the hash symbol ('#') is ignored and can be used to introduce comments or to prevent the given analysis from loading temporarily.

Any non-numeric items found in a data column with one of the names listed in the above dictionary are replaced by 'NA' automatically. At the next stage all fully numeric data columns are stored in a numeric data matrix *WR*. Decimal commas, if present in text file, are converted to decimal points. For any missing major- and minor-element data (SiO2, TiO2, Al2O3, Fe2O3, FeO, MnO, MgO, CaO, Na2O, K2O, H2O.PLUS, CO2, P2O5, F, S), an empty (NA) column is created automatically.

The remaining, that is all at least partly textual data columns are transferred into a data frame *labels*, in which are also stored codes of plotting symbols, their colours and sizes.

Subsequently, several extra parameters are calculated (if not provided by the user already), namely concentrations of some elements from oxides (P, K, Ti, Ba, Sr) in ppm, total iron expressed as ferrous oxide (FeOt), Shand's indexes in molar % (A/NK, A/CNK):

$$A / NK = \frac{Al_2O_3}{Na_2O + K_2O} [mol. \%]$$

$$A / CNK = \frac{Al_2O_3}{CaO + Na_2O + K_2O} [mol. \%]$$

two variants of Mg numbers:

$$mg \# = 100 \frac{MgO}{FeO + MgO} [mol. \%]$$

$$Mg \# = 100 \frac{MgO}{FeOt + MgO} [mol. \%]$$

and K2O/Na2O ratios by weight.

At the same time are the analyses recast to millications (data matrix *milli*) and major-element analyses to anhydrous basis (data matrix *WRanh*).

Data editing and grouping

The *GCDkit* contains basic functions for data input and output, such as pasting, loading and saving or union of two data files (either appending new samples or new data columns for samples already present in the system). If need be, the data set can be modified in a simple spreadsheet. Subsets of the data can be displayed/selected, data columns deleted or new ones appended. In addition, there is a possibility of editing a single value (in the R jargon a *level*) of the given label (a *factor*).

Important feature of data handling in the *GCDkit* is **grouping**. Each of the samples can be assigned to one of the groups that are subsequently utilised by some statistical and plotting functions. Groups can be defined on the basis any of the labels (locality, rock type...), a value of a chosen numerical variable, position in classification diagram (e.g., TAS) or cluster analysis. The information concerning current grouping is stored in a vector *groups*; default grouping after loading a new data file or selection of subset is on plotting symbol.

Searching and subsetting in GCDkit

Great emphasis in the *GCDkit* is on data management, esp. **searching and selecting subsets**.

1. **Searching** is useful for temporary selections of samples that are, for instance, to be displayed or plotted in a diagram. In this case, the data stored in memory are fully unaffected. Search dialog appears automatically within each relevant function (plots, norms etc.)
2. Upon **selection of a subset** (via menu items *Select subset by sample name or label*, *Select subset by range*, *Select subset by Boolean*), the data in memory are replaced by their part (subset) fulfilling the given criteria. Still, the changes are temporary as each time a file is loaded, a backup of the data is kept that can be restored anytime using the menu *Select the whole data set* (connected to the function *selectAll*). However, in that case all changes made e.g. to plotting symbols, grouping etc. would be lost.

The searching and subsetting functions employ powerful regular expressions (see R help for *regex*). Syntax of search patterns employed in searching and subsetting is dealt with in a considerable detail in the corresponding manual/help entries, here are shown only examples.

a) By sample name or label

These queries of a single textual column employ regular expressions [2]. Put in simple terms, most characters, including all letters and digits, are regular expressions that match themselves. However, metacharacters with a special meaning ('?', '+', '{', '|', '(', and ')') must be preceded by a backslash. Two regular expressions may be joined by the infix operator '|'; the result then matches either of them (logical **or**).

| Regular expression | Matches |
|--------------------|-----------------------------------------------------------|
| . | Any character except NULL |
| ^ | Beginning of the expression |
| \$ | End of the expression |
| [] | Any of the characters given in square brackets |
| [m-n] | Any character in the range given by <i>m</i> and <i>n</i> |

A regular expression may be followed by one of several repetition operators.

| Repetition operator | The preceding item will be matched |
|---------------------|-----------------------------------------------------------|
| ? | At most once (i.e. is optional) |
| * | Zero or more times |
| + | One or more times |
| {n} | Exactly <i>n</i> times |
| {n,} | At least <i>n</i> times |
| {n,m} | At least <i>n</i> times, but not more than <i>m</i> times |

As an example, let us consider that the searched field corresponds to the following localities: Mull, Rhum, Skye, Coll, Colonsay, Hoy, Westray, Sanday, Stronsay, Tiree, Islay.

| Search pattern | Matches |
|----------------|-----------------------------|
| ol | Coll, Colonsay |
| n.a | Colonsay, Sanday, Stronsay |
| ^S | Skye, Sanday, Stronsay |
| e\$ | Skye, Tiree |
| [ds]ay | Colonsay, Sanday, Stronsay |
| [p-s]ay | Colonsay, Westray, Stronsay |
| ol oy | Coll, Colonsay, Hoy |
| l{2} | Mull, Coll |

If the sample names are: B1-1, B1-3, Koz-1, Koz-2, Koz-5, Koz-11, KozD-1, Ri-1:

| Search pattern | Matches |
|----------------|-----------------------------------------------|
| oz-[1-3] | Koz-1, Koz-2, Koz-11 |
| oz- B1- | B1-1, B1-2, B1-3, Koz-1, Koz-2, Koz-5, Koz-11 |

b) By range

In this case the search pattern is treated as a selection of sample sequence numbers (effectively a list separated by commas that may also contain ranges expressed by colons). See some examples below:

| Search pattern | Matches |
|----------------|----------------------------------------|
| 1:5 | First to fifth samples in the data set |
| 1,10 | First and tenth |
| 1:5, 10:11, 25 | Samples number 1, 2, ... 5, 10, 11, 25 |

3. By Boolean conditions

The search pattern, representing a Boolean condition, may contain names of both the textual and numeric columns as well as comparison operators '<' (lower than), '>' (greater than), '<=' (lower or equal to), '>=' (greater or equal to), '=' or '==' (equal to), '!=' (not equal to). The character strings should be quoted. Two or more conditions can be combined together by logical **and** (expressed as '.and.', '.AND.', '&'), **or** ('.or.', '.OR.', '|') and brackets.

Example – Subset by Boolean conditions

| Search pattern | Matches |
|---------------------------------------------------------------------------------------|----------------------------------------------------------------------------------------------------|
| Locality="Rhum" | All analyses from Rhum |
| Locality="Rhum".and.SiO2>65 Locality="Rhum".AND.SiO2>65 Locality="Rhum"&SiO2>65 | Analyses from Rhum with SiO ₂ greater than 65 (all three expressions are equivalent) |
| MgO>10&(Locality="Skye" Locality="Islay") | Analyses from Skye or Islay with MgO greater than ten |
| MgO>=10&Locality!="Skye"&Locality!="Islay" | Analyses from anywhere except Skye and Islay with MgO at least ten |

Using multiple datasets

As a relatively new feature, *GCDkit* has introduced a possibility of keeping several datasets in memory together. Only one among them at the time is designated as the so-called current dataset (that one whose name is indicated in the title bar of the Windows R Console Window)

and used for calculations and plotting. The current dataset can be chosen from the menu *GCDkit|Switch/restore a dataset* (attached to the function `selectDataset`) or directly by the function `peekDataset`.

Any new dataset, as soon as it is loaded from file, clipboard or Internet connection, is backed up in the memory under its root filename. If needed, a time stamp is attached to make this name unique.

Modified datasets can be stored/replaced anytime from the menu *GCDkit|Store the current dataset* (function `pokeDataset`).

All the datasets stored in memory, apart from the current one, can be deleted at once (menu *GCDkit|Purge stored datasets*).

Moreover, any of the datasets stored in memory can be added to the current standard (Figaro-compatible) binary/ternary plot or spiderplot, or their plate. This is done either in the foreground or in the background (menus *GCDkit|Overplot another dataset* and *GCDkit|Underplot another dataset*, respectively). These tools allow effortless comparisons, say with real-world data from a similar pluton or a numeric matrix spanning from petrogenetic modelling.

Choosing numeric variable(s), calculation core routine

Many functions (e.g. binary plots, some statistics) require a single numeric variable to be chosen first. The easiest way is to type in the name of the numerical column (e.g., 'SiO2') or its sequence number (2 for the second column). However, it is not necessary to enter the name in its entirety. Only a substring that appears somewhere in the column name or other forms of regular expressions can be specified.

If the result is ambiguous, the correct variable has to be picked manually from a drop-down list of the multiple matches. Ultimately, empty response invokes list of all available variables.

As an useful alternative in many instances, the system allows to enter a formula that can involve any combination of names of existing numerical columns with constants, brackets, arithmetic operators `+-*/^` and R functions. The most useful of these are '*sqr*t' (square root), '*log*' (natural logarithm), '*log10*' (common logarithm) and '*exp*' (exponential function). Potentially useful can be also '*min*', '*max*' (extremes), '*length*' (number of elements/cases), '*sum*', '*mean*', and '*prod*' (product of the elements). In any case, no one prevents the advanced users from writing their own functions that can be invoked here.

When multiple columns are to be selected (e.g. for multiple or correlation plots), the easiest way is to type in directly their names separated by commas. Alternatively can be used a comma-delimited list of sequence numbers that may also contain ranges expressed by colons. Also user-defined or built-in lists can be employed, such as 'LILE', 'REE', 'major' and 'HFSE' or their combinations with the column names. Again, if the input field is left empty, and Enter pressed, a complete list of variables is shown.

Calculations

All the calculation algorithms produce output temporarily stored in the memory, in the variable (a vector, matrix or list) *results*. Before it is rewritten by the next calculation procedure, it can be saved or appended to the current dataset for further processing (e.g., CIPW normative albite, quartz and orthoclase to be plotted in the Ab–Q–Or ternary).

Most of the normative recalculation algorithms have been adopted from an earlier QuickBasic program NORMAN (Janoušek, 2001). Currently are available R modules for calculation of the CIPW norm, including the modification with biotite and hornblende (Hutchison, 1974; 1975), Niggli Catanorm (Hutchison, 1974 and references therein), Niggli's cationic values (Niggli, 1948), multicationic parameters of the French authors (De La Roche

et al., 1980; Debon and Le Fort, 1983; 1988; Bonin *et al.*, 2020), improved Mesonorm for granitoid rocks (Mielke & Winkler, 1979). As an optional outcome of the latter, implemented is a Q'–ANOR binary chemically approximating the modal QAPF classification for the igneous rocks (Streckeisen & Le Maitre, 1979).

The *GCDkit* provides a menu-driven interface to (a fraction of) statistical functions built in R. These include simple descriptive statistics, histograms and boxplots (for the whole data set or individual groups), correlation plots as well as more sophisticated methods of multivariate statistics (cluster analysis and principal components).

Plotting

The plotting symbols and colours can be allocated, if need be independently of each other, according to current grouping or any of the labels or, for colours, numeric variables. If two distinct criteria for symbols and colours are chosen, two legends are built (see <http://blog.gcdkit.org/2017/04/legends.html>).

The *GCDkit* produces publication quality plots in PDF or PostScript formats that can be easily imported into other graphical or DTP package for further editing (simply right-click the graphical window and select the desired format). All the plots can be saved simultaneously to PDF (*Save all graphics to PDF*) or PostScript (*Save all graphics to PS*).

The strength of *GCDkit* consists in the wealth of built-in plots most of which (the stand alone, i.e. not multiple, diagrams) are defined as templates compatible with *Figaro*, a set of graphical utilities for R.

Figaro provides means to create figure objects, which contain both the data and methods to make changes to the figure (via the menu *Plot editing*). So, for example, the title can be changed or the histogram fill colour altered and any changes are automatically made visible on interactive devices. In addition, one can zoom in and out of the data. *Figaro* objects currently permit the editing of the text, font, size and colour of the main title, sub title and axis labels; colour, size, and symbol of points; colour, line type, and width of lines. Thus *Figaro* provides a degree of interactive editing before committing to hardcopy. Many *Figaro* templates are defined as a set of closed polygons; if so, they can be used by the generic classification algorithm.

GCDkit employs the concept of the so-called plates, i.e. collections of two or more *Figaro* plots in a single plotting window. The examples are Harker plots, or diagrams using the same *Figaro* template plotted for individual data groups (menu item “*Multiple plots by groups*” – function *figMulti*), based e.g. on the locality they come from. Alternatively, plates can be fully user-defined and contain any mixture of *Figaro*-compatible plots, such as binary plots, ternary plots, classification diagrams or spiderplots (including selected panels from other plates).

On the Windows GUI, a single plot can be selected by graphical cursor from the plate already plotted (just right click the plot window and invoke the menu option *Select slot – Select another slot*) and subsequently edited in the manner previously available solely to stand-alone *Figaro*-compatible plots (submenu of the same menu called *Plot editing*). Moreover, properties of the whole plate or all its diagrams can be changed simultaneously (submenu *Plate editing*). For instance, the whole plate can be set to black and white, the font size of axis labels can be altered, scaling of the common axis on Harker plots set, or the minima of all y axes of binary plots set to 0.

NOTE Use menu item *Plate redraw* (attached to the function *plateRedraw*), or dedicated *Export to PS (Colour)* and *Export to PS (BW)* menu items from *Plate editing* submenu to produce “clean” version of your plate before exporting, if prior further editing in a specialized graphical program (CorelDraw, Adobe Illustrator...).

Known main differences of implementation on Mac and Linux from the (native) Windows version

- Supported are only reading/writing of text files, clipboard, CSV and DBF files,
- Missing import/export from/to XLS, XLSX and MDB,
- No interactive editing of plates,
- Language versions for classification plots not implemented,
- Command history does not work correctly.

Where to go next?

Arguably a useful introduction to using R with, or without, *GCDkit* in interpretation and modelling of whole-rock geochemical data provides our monograph (Janoušek *et al.*, 2016). Courtesy of the publisher, the Appendices explaining the R syntax and the workings of the *GCDkit* system, are freely available online at: <http://blog.gcdkit.org/2015/10>. But if you want to support our efforts, and learn more, buy a paper or electronic copy!

Future

Our long-term aim is to build a single, coherent and platform-independent system with high-level plotting capabilities for interpretation of whole-rock geochemical data that would be straightforward to use by ordinary users but, at the same time, easily expandable by the more demanding ones. In a nearest future *GCDkit* should be expanded by functions for direct and inverse modelling of the main petrogenetic processes (fractional crystallization, AFC, binary mixing). These are ready but not yet stable enough to be released.

Note that functions for recalculation of mineral chemical data are available already, forming a part of the specialized R package named *GCDkit.Mineral* (Janoušek *et al.* in print), downloadable from <https://mineral.gcdkit.org>.

Enjoy! On behalf of the authors

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<http://www.gcdkit.org>

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