UThwigl - an R package for closed- and open-system uranium-thorium dating

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Abstract

For several decades, uranium-thorium (U-Th) dating has allowed geochronologists to precisely date geological materials, providing invaluable geochronological constraints on Quaternary processes. Open-system dating of bones and teeth has also provided ages of human and faunal remains of archaeological significance.

To facilitate access to closed- and open-system U-Th dating to the broad scientific community, here we provide an R package, named *UThwigl*. Description of input and output parameters is given, as well as a guide for running the model. The package can be used three different ways: (i) as a web application, (ii) through a web browser with an internet connection, or (iii) in R (most efficiently with RStudio). Examples of application of the model are also provided, showing that it yields ages within error of previously published values.

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Introduction

Uranium-thorium (U-Th) dating has revolutionised Quaternary science and archaeology. Dating uses the decay of ²³⁸U into ²³⁰Th, with ²³⁴U and a few short-lived nuclides as intermediary products. It is based on the principle that the age of formation of a material can be dated as it incorporates U and no or little Th at the time of formation, so all the ²³⁰Th in the sample comes from decay of ²³⁸U. If detrital Th is incorporated into the sample, a correction must be included to account for the fraction of ²³⁰Th which is detrital and not derived from ²³⁸U decay. Another requirement is that there is no gain or loss of ²³⁰Th, ²³⁴U or ²³⁸U after formation of the material (closed system).

Closed-system U-Th dating can be used to date materials as young as a few years up to samples over 600,000 years old (Edwards et al., 2003). It has been successfully applied to a range of marine and lacustrine carbonates. For instance, dating of corals (coralline aragonite) has had a pivotal role in reconstructing past sea levels (Lambeck and Chappell, 2001). Closed-system U-Th dating has also been applied to speleothems (cave carbonates) which are commonly used as continental palaeo-climate archives (Richards and Dorale, 2003). In corals and most speleothems, detrital correction is minimal; however, it can be significant when dating pedogenic carbonates, for instance (Ludwig and Paces, 2002). In this case, detrital correction can be performed using the measured or assumed composition of the detrital fraction (e.g. Ludwig, 2003a). Alternatively, isochron techniques can be applied (Ludwig and Titterington, 1994); the latter are beyond the scope of this article but IsoPlot is a commonly used software for isochron calculations and other geochronological applications (Ludwig, 2003b), now also available as a R package (Vermeesch, 2018).

Closed-system conditions are seldom met in shells, teeth and bones (although enamel can sometimes be quite impervious to isotope gain or loss). Thus, U-Th dating requires to take into account open system behaviour. The diffusion-adsorption model of Pike and Hedges (2002) and the Diffusion-Adsorption-Decay (DAD) model of Sambridge et al. (2012) were instrumental to implement successfully open-system U-Th dating. They allow for advective and diffusive transport of uranium and thorium isotopes, while including synchronous radioactive decay. Software implementation for the DAD model was written in Fortran and is available as a Java GUI (http://www.iearth.org.au/codes/iDaD/).

Open-system U-Th dating of teeth and bones, while challenging, has provided quantitative ages for human and faunal remains (Eggins et al., 2005; Grün et al., 2014; Hoffmann et al., 2018; Pike and Hedges, 2002; Sambridge et al., 2012). Thus, this approach has significantly improved our understanding of human evolution (e.g. Dirks et al., 2017; Hoffmann et al., 2018; Sutikna et al., 2016).

In this article, we present a R package, *UThwigl*, which offers functions to perform closed-system, csUTh(), and open-system, osUTh(), U-Th age calculations. The former implements formulations given in Ludwig (2003a) while the latter applies the DAD model of Sambridge et al. (2012). The R package *IsoPlotR* provides a more extensive tool for closed-system U-Th dating (Ver-

meesch, 2018), and *UThwigl* only includes closed-system U-Th age calculations for the sake of offering both closed- and open-system calculations.

Providing an R package aims at increasing the transparency, reproducibility, 48 and flexibility of the analytical workflow for computing U-Th ages. For instance, with open-system dating, it is difficult to include the Java GUI in a 50 fully scripted data analysis so the method for computing the DAD model is not 51 fully transparent. This can obscure steps where key decisions are made that are 52 important for others to see to verify the reliability of the analysis. Enabling a 53 scripted workflow for computational analysis of geoscience data is important for 54 improving the reproducibility of results. Reproducibility refers to the ability to recreate the results or retest the hypotheses leading to a scientific claim, either by rerunning the same code used by the original authors, or by writing new code. 57 High rates of irreproducibility of research results have been estimated in several fields and disciplines (Camerer et al., 2018; Camerer et al., 2016; Freedman et 59 al., 2015; Institute, 2013; Ioannidis, 2005; Open Science Collaboration, 2015). Consequently, the transparency, openness, and reproducibility of results and 61 methods are receiving increased attention, and the norms of research in many 62 fields are changing (Marwick, 2016; Miguel et al., 2014; Nosek et al., 2015). 63

There is strong interest in open, transparent, and reusable research in the geoscience community (Gil et al., 2016) and substantial progress toward open data has been made in the geosciences with the widespread use of data services of NASA, USGS, NOAA and community-built data portals such as OneGeology, EarthChem, RRUFF, PANGAEA, PaleoBioDB, SISAL and others (Comas-Bru et al., 2020; Kattge et al., 2014; Ma, 2018). However, the use of open source software such as R (Pebesma et al., 2012), and sharing of scripted data analysis workflows with research publications is not yet widespread (Hutton et al., 2016). With this R package our goal is to make scripted and reproducible data analysis easy for uranium-thorium dating. This will improve the transparency of geochronology research, and provide a more credible and robust foundation for scientific advancement (Hutton et al., 2016).

To enable re-use of our materials and improve reproducibility and transparency, all the results and visualisations in this paper can be reproduced using the RMarkdown vignette document included with the UThwigl package. We have archived these files at http://doi.org/10.17605/OSF.IO/D5P7S to ensure long-term accessibility. Our code is released under the MIT licence, our data as CC-0, and our figures as CC-BY, to enable maximum re-use (for more details, see Marwick, 2016).

Methods

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Closed-system U-Th dating is based on the premise that the dated material takes up U during formation, but no Th (and thus no ²³⁰Th). This is because U-Th dating focuses on materials that precipitate from solution (e.g. carbonates) and Th has a very low solubility in most cases. In this case, the age of material formation (e.g., precipitation of coralline aragonite) is quantified. Carbonates

are particularly amenable to U-Th dating because, in most cases, they take up U during formation but very little Th.

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U-Th dating is undertaken by measuring the $^{234}\mathrm{U}/^{238}\mathrm{U}$ and $^{230}\mathrm{Th}/^{238}\mathrm{U}$ of the material. In this case, the measured $[^{230}_{238}\underline{U}]$ activity ratio in the sample is written as follows (brackets denote activity ratios):

$$\left[\frac{^{230}Th}{^{238}U}\right] = 1 - e^{-\lambda_{230}t} + \left(\frac{\delta^{234}U_m}{1000}\right) \cdot \left(\frac{\lambda_{230}}{\lambda_{230} - \lambda_{234}}\right) \cdot \left(1 - e^{-(\lambda_{230} - \lambda_{234})t}\right)$$

where λ_{230} and λ_{234} are ²³⁰Th and ²³⁴U decay constants, respectively (in yr⁻¹), t is the age of the material (i.e. the time elapsed since onset of ²³⁰Th in-growth; in yr), and $\delta^{234}U_m = (\left[\frac{^{234}U}{^{238}U}\right] - 1).1000$, with the measured $\left[\frac{^{234}U}{^{238}U}\right]$ activity ratio in the material.

This approach assumes that (i) all the ²³⁰Th measured is produced by decay of ²³⁸U and (ii) the system is closed at t=0, i.e. there is no loss or gain of any nuclides after the time of formation. In corals, the second assumption can be tested using the mineralogy of the sample: corals precipitate as aragonite. Open system behaviour exemplified as diagenetic alteration generally involves the replacement of aragonite by calcite. Thus, prior to ²³⁰Th dating, coral mineralogy should be quantified. Samples exhibiting calcite are deemed unsuitable for dating. The initial $\binom{234}{238U}$ activity ratio, calculated along the age, should also be similar to the seawater value [1.145; Henderson (2002)], which has not changed significantly over the past 800,000 yr. In speleothems, the closed system assumption can be tested by looking for any age inversions. In pedogenic carbonates, there is no straightforward way to test this assumption. The first assumption (²³⁰Th measured in only produced by decay of ²³⁸U) is tested for any sample type using ²³²Th as an index of detrital Th. Any significant amount of ²³²Th in the sample implies a detrital contribution of Th to the sample, and thus that a fraction of the 230 Th measured does not result from decay of 238 U, but from detrital input of Th. The $[^{230}_{\overline{L}h}]$ or $[^{232}_{\overline{L}h}]$ activity ratios are used as indices of the quantity of detrital Th present in the sample. Arbitrary values are set to define whether the presence of detrital Th significant; $\begin{bmatrix} \frac{230}{232}Th}{232}Th \end{bmatrix}$ ratios greater than 20 or $\left[\frac{^{23}Th}{^{238}U}\right]$ ratios less than 0.01 are usually recommended. If significant detrital Th is present, correction is necessary (in fact, even if the contribution of detrital Th is minimal, correction should still be applied). Ideally, one would measure the $\left[\frac{230}{238}\frac{Th}{U}\right]$, $\left[\frac{234}{238}\frac{U}{U}\right]$ and $\left[\frac{230}{232}\frac{Th}{Th}\right]$ activity ratios of the detrital component; however it is rarely possible to isolate this fraction, let alone measure its U-series isotope composition. Often, correction is undertaken assuming a $\left[\frac{230}{232}\frac{Th}{Th}\right]$ of 0.8 \pm 0.4 for the detrital component, which is the average value for the continental crust. Alternatively, detrital correction can be undertaken by measuring several samples assumed to have the same ²³⁰Th age, but variable amounts of detrital Th. In this case, it is possible to define an isochron or derive a single age for the same of isochronous samples.

Open-system U-Th dating operates on the principles that little U (and Th) are incorporated at the time of the material formation (shell, tooth or bone),

and it is only after death and burial of the material in soil or sediments, that U (and Th to a lesser extent) are taken up and diffuse into the material. When dating teeth, enamel is often preferred over dentine, as the former is denser and thus less prone to complex nuclide movement. For samples exhibiting open system behaviour, the analytical strategy generally involves measuring $^{238}\mathrm{U},^{234}\mathrm{U},^{230}\mathrm{Th}$ and $^{232}\mathrm{Th}$ in several aliquots along a transect perpendicular to the surface of the sample (Figure @ref(fig:femurpic)). Then, U concentrations and $[\frac{^{230}Th}{^{238}U}],~[\frac{^{234}U}{^{238}U}]$ activity ratios can be modelled to derive a single open-system age. Aliquots can be collected by micro-drilling or using laser ablation.

Several open-system models have been developed (Pike and Pettitt, 2003). The Diffusion-Adsorption model (Pike and Pettitt, 2003) was later modified to a Diffusion-Adsorption-Decay model (Sambridge et al., 2012), and is the most commonly employed to U-Th date archaeological materials such as teeth and bones. Profiling of uranium concentrations across the sample is used to determine whether the sample has experienced loss of uranium (inverted "U" shaped profile) or shows an irregular pattern of uranium concentration variation. If the sample presents either of these profiles, it is rejected for dating. Ideally, the uranium concentration profile shows a "U" shape (illustrating uranium diffusion) or homogenous concentrations (indicating that equilibrium in uranium diffusion has been achieved). Once these tests have been performed, closed-system U-Th ages for each analysis can then be computed. If they show an inverted "U" shaped profile, this is diagnostic of recent uranium uptake, and the sample is rejected. Otherwise, the profile of U-series isotope data can then be used to derive a single open-system age.

Analytically, two types of measurements are possible: bulk or in-situ. For bulk analysis, a fraction of the samples is dissolved and the solution processed through ion exchange chromatography to separate U and Th (e.g. Luo et al., 1997). Each element is then analysed separately for their isotope ratios by mass spectrometry. For in-situ analysis, laser ablation is commonly used (Eggins et al., 2005). In this case, a laser with a spot size ranging from a few μm to several hundreds of μm produces an aerosol which is carried using a gas (helium or preferably a mixture of helium and nitrogen; Eggins et al. (1998)). While laser ablation offers a better spatial resolution and is less time consuming than bulk analysis, the precision of the data is inferior because of the much smaller amount of material sampled.

Uranium and thorium isotope ratios are analysed by multi-collector inductively-coupled plasma mass spectrometry (e.g. Luo et al. (1997); although bulk analysis can also be performed by thermal ionisation mass spectrometry). A plasma ionises U and Th atoms, their isotopes are separated through a magnetic field and each are collected in a different collector (Faraday cups or ion counters). If using laser ablation, it is best to have two ion counters so ²³⁰Th and ²³⁴U can be collected simultaneously.

Closed-system dating

Pending closed-system behaviour can be assessed, it is possible to derive an age for each U-Th analysis. The closed-system function csUTh() requires that

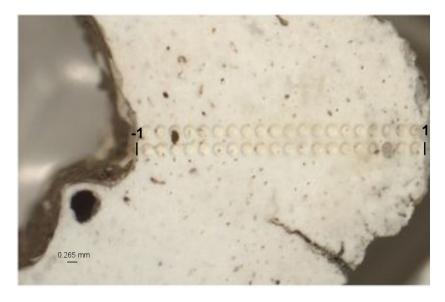


Figure 1: (ref:femurpiccap)

(#fig:femurpic)

for each analysis to yield an age, $\left[\frac{234}{238U}\right]$, $\left[\frac{230}{238U}\right]$ are measured, as well as $\left[\frac{232}{238U}\right]$ or $\left[\frac{230}{232Th}\right]$. The $\left[\frac{232}{238U}\right]$ activity ratio is required for detrital correction (note it is needed to use csUTh() whether the detrital correction is performed or not). If $\left[\frac{230}{232Th}\right]$ is provided instead, $\left[\frac{232}{238U}\right]$ is calculated with csUth().

Open-system dating

Data required for the DAD model are $\left[\frac{^{230}Th}{^{238}U}\right]$ and $\left[\frac{^{234}U}{^{238}U}\right]$ activity ratios collected along a transect perpendicular to the surface of the tooth or bone.

The x-y coordinates of each analysis, and of the inner and outer surfaces of the sample are also needed as input data. osUTh() uses these coordinates to calculate normalised positions, where the outer surface of the sample is given a reference coordinate of 1, the inner surface -1, and analyses take values in between (Figure @ref(fig:femurpic)).

(ref:femurpiccap) Modern human femur (132A/LB/27D/03) from Liang Bua, Flores, Indonesia. Two analysis transects can be seen. For a given transect, the x and y coordinates of the outer and inner surfaces, and of the analyses, are used by osUTh() to calculate normalised positions where the outer surface is given 1 as reference coordinate, the inner surface -1, and normalised positions of the analyses take values in between. Modified from Sutikna et al. (2016).

Working with the package

We provide three methods for using this package to suit different levels of familiarity with the R programming language. The simplest way to use the pack-

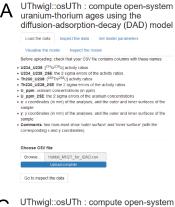
age is our web applications, online at https://anthony-dosseto.shinyapps.io/csUTh/ and https://anthony-dosseto.shinyapps.io/osUTh/ (Figure @ref(fig:shinyfig)). Using the web application requires no familiarity with R. To use the web application we upload a CSV file, then click through a series of tabs to inspect the data, adjust the model parameters, run the model, and inspect the output. The interface is mouse-driven and requires no programming. In the web application we upload the data file on the Load the data tab, set parameters from the Set model parameters tab, run the model by clicking the button Run Simulation on the same tab, and observe the results on the Visualise the model and Inspect the model tabs. We can change the parameters and re-run the model by click the button Run Simulation. Once done, close the window.

(ref:shinyfigcap) Screenshots of the web application for open-system U-Th dating. A: Upload a CSV file of the data to model, B: Inspect a table of the uploaded data. C: Set the model parameters and run the model. D: Inspect visualisations of the model's output. E: Inspect and download the numeric output from the model.

The second way to use the package is with Binder, a browser-based instance of R and RStudio that includes our package ready to work with (Figure @ref(fig:binderfig)). Binder is a server technology that turns computational material, such as an R package, into interactive computational environments in the cloud. Using Binder requires a novice level of familiarity with R, for example to use the code in this paper and adapt it to work with a different CSV file. Because Binder provides a complete R environment, custom R code can be written during a Binder instance to further explore the model's output in the browser. These two methods, the web application and Binder, do not require any software to be downloaded and installed on the user's computer, all computation occurs in the browser. The web application and Binder are suitable for getting a quick start on working with the package, but they require a connection to the internet, and they have limited memory and compute time available per instance.

(ref:binderfigcap) Screenshot of Binder running R and RStudio in a web browser window.

The third method is to download and install the package locally to the user's computer, and work with it in the user's local installation of R and RStudio. This method requires some familiarity with R, but gives the most flexibility when working with the model because we are not limited by the memory and compute time of the cloud services. Our recommendation is to use Binder or a local installation of UThwigl because then the user can save an R script file that includes the name of the input file, the specific parameters used to generate the model output, and any downstream processing and visualisation. This script file and the CSV file can then be archived in a data repository to ensure long-term accessibility for other researchers. In the following sections we demonstrate the use of UThwigl with a local installation of R and RStudio.



C UThwigl::osUTh : compute open-system uranium-thorium ages using the diffusion-adsorption-decay (DAD) model



UThwigl::osUTh : compute open-system uranium-thorium ages using the diffusion-adsorption-decay (DAD) model



B UThwigl::osUTh: compute open-system uranium-thorium ages using the diffusion-adsorption-decay (DAD) model



UThwigl::osUTh : compute open-system uranium-thorium ages using the diffusion-adsorption-decay (DAD) model

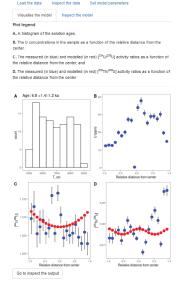


Figure 2: (ref:shinyfigcap)

(#fig:shinyfig)

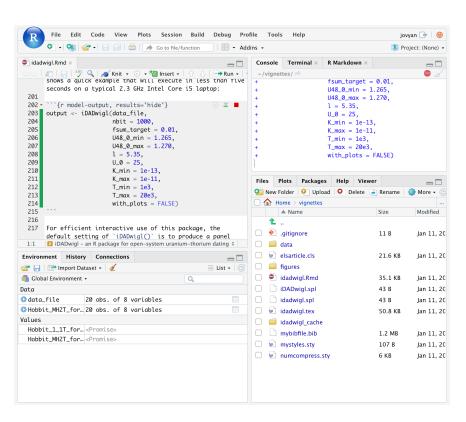


Figure 3: (ref:binderfigcap)

(#fig:binderfig)

Installing and attaching the package

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First the user will need to download and install R, and we also recommend downloading and installing RStudio. To run the model, start RStudio and install the package from GitHub. There are many ways to do this, one simple method is shown in the line below. This only needs to be done once per computer.

```
if(!require("remotes")) install.packages("remotes")
remotes::install_github("tonydoss/UThwigl")
```

For routine data analysis, R scripts need to contain the following line to attach the package to the current working environment. This line needs to be run at the start of each analysis:

```
# attach the package
library(UThwigl)
```

Closed-system U-Th dating

Input data format 249

> Our package provides the fuction csUTh() for closed-system U-Th dating. Data for this function needs to be in a data frame (a form of table in R) with the following column names:

- Sample_ID 253
 - U234_U238
 - U234_U238_2SE
 - Th230_U238
 - Th230_U238_2SE

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- Th232_U238
 - Th232_U238_2SE

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- Th230_Th232
- Th230_Th232_2SE

To help with preparing data for input into our function, we have included an example of an input file, taken from Pan et al. (2018). Before reading in the data file, the user needs to set the working directory to the folder containing the data file. This can be done in RStudio using the menu item 'Session' > 'Set Working Directory' > 'To Source File Location.' Alternatively, the working directory can be defined interactively at the R prompt in the Console panel useing setwd(). However, we do not recommend including setwd() in script files because it is bad for reproducibility, since the path to one user's working directory will not exist on another user's computer.

Inspecting the included data sets will be helpful for understanding how to prepare new data for use with this package. After attaching the package, we can access the built-in datasets with the data() function, like this: 275

access the data included in the UThwigl package data("Pan2018")

This will make the built-in data available in the R environment to inspect and explore how to use the csUTh() function. To download the built-in data to the user's computer as a CSV file, so it can be inspected and modified in a spreadsheet program (e.g. as a template for the user's own data), use write.csv():

```
# download the data included in the package
write.csv(Pan2018, "Pan2018.csv")
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The code chunk below shows how to read the CSV file created above into the R environment. We assume that the user's working directory contains a directory called data and the CSV file is in this data directory, and so the data can be imported as follows:

```
# read in one of the example CSV files included in the package
input_data_cs <- read.csv('data/Pan2018.csv')</pre>
```

To use new data with this package, the user needs to import a CSV or Excel file with the U-Th data into the R environment. This can be done using a generic function such as read.csv or read_excel from the readxl package (Wickham and Bryan, 2018).

Table @ref(tab:pan) shows the data contained in the Pan2018.csv file included in the package.

		2SE		2SE		2SE
	U238	U238_	_U238	_U238_	_U238	_U238_
${ m Sample}_{ar{}}$	0.234_1	0234_1	${ m Th}230$	${ m Th}230$	${ m Th}232$	${ m Th232}_{_}$
$\overline{\mathbf{S}_{a}}$	î	ŢŊ	Ī	Ē	Ē	Ē
YP002A	1.150	0.005	0.794	0.007	0.010	0.00005
YP002B	1.120	0.004	0.788	0.006	0.004	0.00002
YP003-1_1	1.125	0.004	0.752	0.010	0.000	0.00001
$YP003-1_2$	1.113	0.007	0.761	0.011	0.000	0.00000
YP003-1_3	1.122	0.005	0.748	0.008	0.001	0.00001
$YP003-1_4$	1.122	0.005	0.726	0.007	0.001	0.00001
$YP003-1_{5}$	1.119	0.006	0.757	0.006	0.002	0.00001
YP002-1_1	1.129	0.006	0.722	0.008	0.001	0.00001
$YP002-1_2$	1.137	0.005	0.767	0.008	0.001	0.00001
$YP002-1_3$	1.118	0.008	0.739	0.009	0.002	0.00002
$YP002-1_4$	1.114	0.006	0.749	0.008	0.003	0.00003
$YP002-1_{5}$	1.105	0.007	0.764	0.011	0.003	0.00004

Table 1: Data contained in the example CSV file Pan2018.csv included in the package

The columns Sample_ID, U234_U238, U234_U238_2SE, Th230_U238, Th230_U238_2SE and either Th232_U238 and Th232_U238_2SE, or Th230_Th232 and Th230_Th232_2SE must be present in the input data frame with these exact names for the model to function. The csUTh() function will check if the input data frame has these columns, and will stop with an error message if it does not find these columns. The names() function can be used to update column names of a data frame to

ensure they match the names that the model function requires. Alternatively the user can edit the column names in a spreadsheet program such as Microsoft Excel. The order of the columns in the data frame is not important.

Columns U234_U238 and U234_U238_2SE are the $\left[\frac{234}{238U}\right]$ activity ratios and their 2σ errors. Columns Th230_U238 and Th230_U238_2SE are the $\left[\frac{230}{238U}\right]$ activity ratios and their 2σ errors. Columns Th232_U238 and Th232_U238_2SE are the $\left[\frac{232}{238U}\right]$ activity ratios and their 2σ errors. Columns Th230_Th232 and Th230_Th232_2SE are the $\left[\frac{230}{232U}\right]$ activity ratios and their 2σ errors.

If Th230_Th232 and Th230_Th232_2SE are provided instead of Th232_U238 and Th232_U238_2SE, columns Th232_U238 and Th232_U238_2SE are calculated by csUTh().

Details of the input parameters of closed-system analysis

sample_name is the name of the sample to calculate closed-system ages for. The function will partially match by sample prefix. For example in Table @ref(tab:pan) one sample is indicated by the Sample ID 'YP003.' If the user inputs 'YP003' for the sample_name, then this will match rows where the Sample ID is 'YP003-1,' 'YP003-2,' 'YP003-3,' and so on.

nbitchoice is the number of iterations in the model (it is recommended to have at least 10,000). detcorrectionchoice is a parameter for choosing whether or not to apply a detrital correction to the calculation.

R28det (0.8) and R28det_err (0.4) are the values for the $\left[\frac{^{232}Th}{^{238}U}\right]$ activity ratio of the detritus and its standard error (default values in parentheses). Similarly, R08det (1) and R08det_err (0.05) are the values for the $\left[\frac{^{230}Th}{^{238}U}\right]$ activity ratio of the detritus and its standard error, and R48det (1) and R48det_err (0.02) are the corresponding values for $\left[\frac{^{234}U}{^{238}U}\right]$ activity ratio of the detritus.

How to run the model

Assuming that the package is attached with library(UThwigl), and the data have been imported to the working environment as noted above into a data frame named input_data_cs, the user can then run csUTh(), specifying the input data frame and the input parameters as described above. The code block below shows a typical example that will execute in less than a minute on a typical 2.3 GHz Intel Core i5 laptop:

```
# Solve for sample YPOO3
output_cs <-
csUTh(
   input_data_cs,
   sample_name = 'YPOO3',
   nbitchoice = 10000,
   detcorrectionchoice = TRUE,
   R28det = 0.8,
   R28det_err = 0.4,</pre>
```

```
R08det = 1,
R08det_err = 0.05,
R48det = 1,
R48det_err = 0.02,
keepfiltereddata = FALSE,
print_age = TRUE,
with_plots = TRUE,
save_plots = FALSE,
save_output = FALSE
)
```

For efficient interactive use of this package, the default setting of csUTh() is to produce a panel plot as seen in Figure @ref(fig:csuthvizfig). The setting with_plots = FALSE prevents plots from being generated which is more useful when the function is part of a longer sequence of code. The function runs faster when not producing plots, which is helpful when replicating many runs. The setting save_output = TRUE will save a csv file to the current working directory so the output data can be used in other contexts. The csv file that is created when save_output = TRUE will be given a name that includes a date and time stamp so that the output of each time the function is run can be saved to a unique file.

When run on the R console, the function will print a confirmation that the input data frame has the required columns. If print_age is set to TRUE, it will also print the resulting mean age value of several analyses on a single sample, with an error reported as 2 Standard Deviation, for example:

```
All required columns are present in the input data
[1] "Mean age: 117.1 +/- 3.7 ka"
```

print_age should be set to FALSE if ages computed are not for analyses of the same sample, since this mean age would be meaningless.

Inspecting and visualizing the models' output

The function returns a data frame with the age, error and summary output for each measurement, as shown in Table @ref(tab:panoutput). This includes calculated ages (with or without detrital correction, depending how detcorrectionchoice was set), initial $\left[\frac{234}{238}U\right]$ activity ratios, along with their uncertainties, calculated as the 2.1 and 97.9 percentiles of the population of solutions determined with the Monte Carlo simulation.

The plots produced by the csUTh() function are stored as list objects in the output of the function. We can show the plots by accessing the list like this:

(ref:csuthvizfigcap) Example of the visualisations produced by the csUTh() function, using the demonstration run described above, and five in-situ analyses by laser ablation of coral sample YP003. A: closed-system ages and B: initial $\begin{bmatrix} 234 U \\ 238 II \end{bmatrix}$ activity ratios for each sample analysis.