

# Hands-on Bayesian age-model software

OxCal  
Bchron  
Bacon

# Today

09:00 – 11:00 OxCal, Bchron, Bacon

11:30-13:30 Analysis multiple sites

14:30 Excursion + dinner

*Tomorrow: discuss potential papers*

# OxCal

Most popular Bayesian  $^{14}\text{C}$  chron builder?

Chron. ordering, outlier analysis

Deposition models

Linking multiple sites

Several ways to enter dates: manual,  
graphical, code

Important: be careful with the code `() "" {} ;`

# OxCal deposition models

file:///C:/Program  
%20Files/OxCal/oxcalhelp/hlp\_analysis\_op  
er.html#deposit

Trees

Uniform

Sequence

P\_Sequence

# Running OxCal

Unzip OxCalDistribution.zip

- To C:\Program Files (Windows)
- To /Applications/ (Mac)

Open Firefox

Open OxCal file in Firefox (ctrl+o):

C:\Program Files\OxCal\Index.html (Windows)

[File:///Applications/OxCal/Index.html](file:///Applications/OxCal/Index.html) (Mac)

Agree to warning

Perhaps bookmark the file

# Running OxCal

Open OxCal.txt in text editor

Copy first lines, `D_Sequence() {...};`

In OxCal, File → New

Change view with text icon

Replace code with the copied code

File → Save...

Run

View → Plot dates

# Running OxCal

Repeat actions for other examples:

In OxCal, File > Open *the previous file*

Replace old code with new code, Run

V\_Sequence() {...}; (wiggle-matching)

U\_Sequence() {...}; (constant deposition)

P\_Sequence(10) {...}; (Poisson model)

P\_Sequence(1) {...}; (more flexible)

# Writing OxCal code

Open MSB2K.dat in text editor (is in Bacon/Cores/MSB2K)

Change code to run a P\_Sequence

Beware of () {} "" ;

Oldest dates first!

# Bchron

Haslett and Parnell 2008 (JRSSC, 57: 399-418)  
Parnell et al 2008 (QSR, 27: 1872-1885)

# BChron

Be connected to the Internet

Open R and type in terminal:

```
install.packages('BChron')
```

Choose a nearby mirror (USA)

```
library('BChron')
```

Type `help(BChron)` and follow install steps

# Installation instructions

1. Create a directory on your hard drive called Bchron.
2. Navigate to the R directory and find the Bchron sub-directory within. On windows this will be C:\ program files\R\R-XXXXX\ library\ BChron\ where XXXXX is the version number of R. On other platforms, this directory can be found by typing `.libPaths()` at the R command prompt.
3. In this directory, there there should be three subdirectories called Input, Output, and CalCurve. Copy these to the Bchron directory you created in step 1.

# Example Bchron run

1. At the command prompt in R, type `library(Bchron)`
2. Type `Bchronmenu()` and choose option 1.
3. Locate your Bchron directory, and select `IntCal09.bch` as your calibration curve and `Glendalough.dat` as your input file. All other options can be left as default.
4. Choose option 2 to calibrate the  $^{14}\text{C}$  dates (standard length).
5. Choose option 3 and 'standard' to run the Bchron model.
6. Choose option 4 to run the prediction stage and create a plot of the data. ....laptop may become hot!!!

# Bacon

Unzip Bacon.zip to somewhere nice

Open R and change dir to where Bacon lives

Load the Bacon R code:

```
source('Bacon.R')
```

Run the default core + settings (MSB2K):

```
Bacon()
```

This will run MSB2K at a default resolution of 5  
cm, sample size 1,000

# Ghost plots

Previous plot has sub-panels, remove:

Close the graph by mouse or type `dev.off()`

`proxy.ghost(1)`

`proxy.ghost(2)` .... what is happening?

Open `MSB2K_proxies.csv` in, e.g., Excel

# Dealing with outliers

Bacon("RLGH3")

Do you agree with the proposed model?

# Longer cores

Type: Cores()

Run a much longer core, at lower resolution:

```
Bacon("LesEchets", 100)
```

How old is the core?

What are those light blue dates?

Greyscale not very visible, re-draw:

```
Bacon.PlotAgeModels(info, dark=200)
```

# WLM

Bacon('WLM19')

Open WLM19.csv in text editor

what are all those columns?

If strange extrapolation warnings: remove or correct  
automatically produced `_priors.txt`

Effect of resolution, e.g., 2, 5, 10?

Effect of accumulation rate prior?

Change memory to, e.g., strength 10, mean 0.2 (or 0.95)

# Events

```
AgesOfEvents(yrmin=0, yrmax=1000,  
             window=100, move=10, info)
```

Check file WLM19\_probs.txt

In case of problems, load and run core anew:

```
E.g.: source("Bacon.R")
```

```
Bacon("WLM21", 10)
```

# Reload existing runs

```
Bacon("MSB2K", run=F)
```

```
Bacon.PlotAgeModels(info)
```

(to get graph settings right)

# Known synchronous events

Open Tephra.txt in text editor

What does this do?

What does the magic **=** do?

Run this in OxCal (takes time)

# Testing for synchronous events

In R, go to the Bacon dir and load Bacon.R

Run the first of three WLM cores, WLM19:

```
Bacon("WLM19")
```

Open /WLM19/WLM19\_events.txt in editor

1's indicate depths with wet-shifts, 0 not

If commas between columns, remove them...

# Testing for synchronous events

In R, find the ages of all these events:

```
AgesOfEvents(yrmin=0, yrmax=1000,  
             window=100, move=10)
```

This will write a file ...WLM19\_21\_probs.txt

# Testing for synchronous events

Do the same for cores WLM20 and WLM21  
(remove commas in `_events.txt` file)

# Testing for synchronous events

Now we have three times timing of events

```
Events19 <- read.table("Cores/WLM19/WLM19_21_probs.txt")
```

```
Events20 <- read.table("Cores/WLM20/WLM20_10_probs.txt")
```

```
Events21 <- read.table("Cores/WLM21/WLM21_14_probs.txt")
```

```
plot(Events19, type="l")
```

```
lines(Events20, col="red")
```

```
lines(Events21, col="blue")
```

# Testing for synchronous events

Prob. of events taking place in the 3 sites  
between yrmin and yrmax:

$$- p(\text{WLM19}) * p(\text{in WLM20}) * p(\text{WLM21})$$

$$\text{pr} \leftarrow \text{Events19[,2]} * \text{Events20[,2]} * \\ \text{Events21[,2]}$$

```
lines(Events19[,1], pr, lwd=3, col="green")
```