### Open Science Improvements in Dendrochronology and Data Processing Pipelines for Quantitative Wood Anatomy

Julie Edwards (she/her) University of Arizona, Laboratory of Tree-Ring Research Nov 29<sup>th</sup> 2023

# **Roots for Resilience (R4R)**



# Arizona Institute for Resilience





RESEARCH, INNOVATION & IMPACT Data Science Institute

# **Roots for Resilience (R4R)**

•Develop data science capabilities across the AIR participating departments and research groups

•Accelerate research projects of participating fellows and their home department research groups

•Build professional networks for addressing large-scale challenges and research questions of interest to AIR faculty

•Develop new interdisciplinary collaborations across AIR, DSI, CyVerse, and other academic units for writing new proposals

•Develop a cohort among participants (and Data Science Ambassadors) to support each other in their own research and efforts to engage their departments



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### Foundational Open Science Skills

CyVerse's 8-week virtual workshop teaches you the principles, practices, and how-tos for doing collaborative open science using cutting-edge, open source cyberinfrastructure, in a collaborative, hands-on setting. To see how our FOSS workshop can support your work, check out the curriculum.

### YEAR OF OPEN SCIENCE



Open Access Publications

Open Data

Open Educational Resources

Open Methodology

Open Peer Review Open Source Software





### Examples of Open Science In Dendrochronology

#### **ITRDB** International Tree-Ring Data Bank



#### Programs Name Function ARSTAN\* Chronology development, plotting and analysis Signal Free Chronology development, plotting and analysis SigFree\* Cofecha<sup>ŧ</sup> Dating and Measurement Quality Control Xdate Date floating series Ind2Col Reformat chronology files to space delimited column files. Reformat ring-width decade files to space delimited files Column Reports the content of increment and chronology files MakeRec

← → C 🏻 dmeko.ltrr.arizona.edu/toolbox.html

### **Tree-Ring Matlab Toolbox**

- Description
- <u>Requirements</u>
- Notation
- Function List

Home Vita Course Toolbox Contact



Advanced open-source tools for paleoenvironmental reconstruction Andy Bunn, Kevin Anchukaitis, Tyson Swetnam, Ed Cook, Ifeoluwa Godwin Ale, and Michele Cosi



AGS - 2054514

Motivation (1)

# Legacy, platform-specific compiled code

#### Advanced techniques

#### Reproducible research





Received 31 March 2006; accepted 18 January 2007



 $\nabla$  Research material easiest to reuse and extend

### Motivation (2)

#### cloud-based analysis

xDateR 1. Introduction and Upload	Welcome to the beta varian of <b>vDetaB</b> , a Shiny and far areadeting tree ring data. This and will help you do	CYVERSE Tools - Services - Learn - Collaborate - About - Search
Upload (drag or browse) a standard file of ring widths (.rwl) readable by read.rwl (e.g., Tucson, Heidelberg, compact, TRiDaS format). Dated Series Browse No file selected Or use example data	welcome to the beta version of <b>xDater</b> – a Sinity app for clossidating tree-ring data. This app will help you do initial quality control on a set of (usually) ring widths and potentially identify dating problems. The idea for this app is to use existing functions in the pacakge <i>dpIR</i> to provide an interactive environment for assessing dating quality and potentially fixing problems. A video showing how to use some of the functions is on YouTube. This app contains tools for crossdating data uploaded in an accepted standard <i>nvl</i> file format. You should start	
	by uploading a file using the widget on the left. If you want to use an example data set you can check the "Or use example data" box. The example data are ring-width data where the dates in some of the series have been deliberately corrupted.	The Open Science Workspace for
	After uploading, standard descriptive statistics and plots are produced in the <b>Describe RWL Data</b> tab. This is the beginning of the cross-dating process. The <b>Correlations Between Series</b> tab produces standard correlation graphs and statistics based on segment length and other options specified by the user. The	Collaborative Data-driven Discovery
	Individual Series Correlations tab allows individual series to be further investigated via running and cross- correlation analyses. Rings can be added and deleted using the <b>Edit Series</b> tab. Reports can be saved at every step and a <i>rwl</i> file can be written.	Create Account
	Preliminary functionality to work with undated series (floaters) is available as well. To access those features, first load a dated (master) <i>rwl</i> file. After that, you can then upload a <i>rwl</i> file with arbitary dates. Once you do so, the <b>Undated Series</b> tab will appear. There you can to try to find dates for each of those series as compared to the dated <i>rwl</i> file. As before, you can use a demo data.	
	This project is on GitHub as is dpIR.	
	Please cite <i>dpIR</i> if you use this app in any published work:	
	<ul> <li>Bunn AG (2008). "A dendrochronology program library in R (dplR)." <i>Dendrochronologia</i>, 26(2), 115-124.</li> <li>ISSN 1125-7865, doi: 10.1016/j.dendro.2008.01.002 (URL: http://doi.org/10.1016/j.dendro.2008.01.002).</li> </ul>	
	<ul> <li>Bunn AG (2010). "Statistical and visual crossdating in R using the dplR library." Dendrochronologia, 28(4), 251-258. ISSN 1125-7865, doi: 10.1016/j.dendro.2009.12.001 (URL: http://doi.org/10.1016/j.dendro.2009.12.001).</li> </ul>	
	This app in still in develpoment. Suggestions are welcome!	PANGEO
		FANGLU

Remember! Users should never rely purely on statistical crossdating but always go back to the wood to verify what is actually happening.

A community platform for Big Data geoscience

-Andy Bunn

# **openDendro** Motivation (3)

#### collaborative open source software development

dplR: Dendrochronology Program Library in R		
Perform tree-ring analyses such as detrending, chronology building, and cross dating. Read and write standard file formats used in dendrochronology.		
Version:	1.7.4	
Depends:	$R (\geq 3.5.0)$	
Imports:	graphics, grDevices, grid, stats, utils, <u>lattice</u> ( $\geq$ 0.13-6), <u>Matrix</u> ( $\geq$ 1.0-3), <u>digest</u> ( $\geq$ 0.2.3), <u>matrixStats</u> ( $\geq$ 0.50.2), <u>png</u> ( $\geq$ 0.1-2), <u>R.utils</u> ( $\geq$ 1.32.1), <u>stringi</u> ( $\geq$ 0.2-3), <u>stringr</u> ( $\geq$ 0.4), <u>XML</u> ( $\geq$ 2.1-0), <u>plyr</u> ( $\geq$ 1.8), <u>signal</u> , <u>boot</u>	
Suggests:	<u>Cairo</u> ( $\geq$ 1.5-0), <u>dichromat</u> ( $\geq$ 1.2-3), <u>foreach</u> , <u>forecast</u> ( $\geq$ 3.6), <u>gmp</u> ( $\geq$ 0.5-5), <u>iterators</u> , <u>knitr</u> , <u>RColorBrewer</u> , <u>rmarkdown</u> , <u>testthat</u> ( $\geq$ 0.8), <u>tikzDevice</u> , <u>waveslim</u>	
Published:	2022-06-23	
Author:	Andy Bunn [aut, cph, cre, trl], Mikko Korpela [aut, cph, trl], Franco Biondi [aut, cph], Filipe Campelo [aut, cph], Pierre Mérian [aut, cph], Fares Qeadan [aut, cph], Christian Zang [aut, cph], Allan Buras [ctb], Alice Cecile [ctb], Manfred Mudelsee [ctb], Michael Schulz [ctb], Stefan Klesse [ctb], David Frank [ctb], Ronald Visser [ctb], Ed Cook [ctb], Kevin Anchukaitis [ctb]	
Maintainer:	Andy Bunn <bunna at="" www.edu=""></bunna>	
License:	<u>GPL-2</u>   <u>GPL-3</u> [expanded from: GPL ( $\geq$ 2)]	

Workflow



#### Bootcamp

Register Now!

openDendro Bootcamp

Jan 8<sup>th</sup>-9<sup>th</sup> 2024

Laboratory of Tree-Ring Research

University of Arizona

#### ≽ Announcement

We are pleased to announce a comprehensive two-day bootcamp focused onopenDendro. This event will provide participants with valuable hands-on training in the R and Python programming languages, using openDendro for analysis in dendrochronology.



### Data Processing Pipelines for Quantitative Wood Anatomy

#### Cell measurements





### **Eidg. Forschungsanstalt WSL** Forschung für Mensch und Umwelt



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Marc Katzenmaier<sup>a,b,\*</sup>, Vivien Sainte Fare Garnot<sup>a</sup>, Jesper Björklund<sup>b,c</sup>, Loïc Schneider<sup>b,c</sup>, Jan Dirk Wegner<sup>a</sup>, Georg von Arx<sup>b,c</sup>

<sup>a</sup> Institute for Computational Science, University of Zurich, Winterthurerstrasse 190, 8057 Zurich, Switzerland

<sup>b</sup> Swiss Federal Institute for Forest, Snow and Landscape Research WSL, Zuercherstrasse 111, 8903 Birmensdorf, Switzerland

<sup>c</sup> Oeschger Centre for Climate Change Research, University of Bern, Hochschulstrasse 4, 3012 Bern, Switzerland





#### **ROXAS** analysis cycle



Workflow



#### $\equiv$ README.md

#### Install

create a python environment and run the following command to install all requirements

pip install torch torchvision opencv-python numpy pytorch-lightning segmentation-models-pytorc 🖵 il

#### Usage

Download the model from here and run the model with

python run\_model.py --input=./input/001.jpg --output=./output/001.png

Q

#### •••

#### 🔁 TowardsRoxasAI-main — -zsh — 140×24

Last login: Sun Nov 26 17:12:38 on ttys000 (base) julieedwards@JuliesMacBookPro ~ % cd Documents/Projects/QWAcode/TowardsRoxasAI-main/input (base) julieedwards@JuliesMacBookPro input % ls 002.jpg 001.jpg TAC03A\_2\_1.jpg (base) julieedwards@JuliesMacBookPro input % cd .. (base) julieedwards@JuliesMacBookPro TowardsRoxasAI-main % ls README.md index.html \_config.yml input assets output final\_semseg\_coniferen\_model.pth run\_model.py (base) julieedwards@JuliesMacBookPro TowardsRoxasAI-main % conda activate ROXASAI (ROXASAI) julieedwards@JuliesMacBookPro TowardsRoxasAI-main % python run\_model.py --input=./input/001.jpg --output=./output/001.png load model run model save output finished successfully

(ROXASAI) julieedwards@JuliesMacBookPro TowardsRoxasAI-main %



### Data Processing Pipelines for Quantitative Wood Anatomy



The University of Arizona is on the land and territories of Indigenous peoples, with Tucson being home to the O'odham and the Yaqui

# Thank you!









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