

Code Ocean eScience presentation

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A really important quote we should all think about

“[A]n article about computational result is advertising, not scholarship. The actual scholarship is the full software environment, code and data, that produced the result.” David Donoho, paraphrasing John Claerbout

Code Ocean

- Code Ocean is, more or less:
 - ▶ JupyterLab IDE + modifications
 - ▶ A robust dependency management system
 - ▶ A publishing platform (DOIs & stable URLs)
 - ▶ A sharing platform (embed your 'compute capsules' on webpages)

The screenshot displays the Code Ocean web interface. At the top, a blue banner indicates the capsule is 'Published' and provides the title 'A Standard for the Scholarly Citation of Archaeological Data' by Ben Marwick et al. Navigation tabs include Capsule, File, Edit, View, Tabs, Settings, and Help. On the right, there are buttons for 'Metadata', 'Edit Your Copy', and a user profile icon.

The left sidebar shows a file explorer with a tree view of the capsule's contents:

- environment (957 B)
- code (100.76 KB)
 - LICENSE (1.04 KB)
 - paper.Rmd (98.73 KB)
 - README.md (881 B)
 - run.sh (133 B)
- data (Manage Datasets) (306.69 MB)
 - figures (2.29 MB)
 - raw_data (304.39 MB)
 - LICENSE (6.4 KB)
 - .gitignore (7 B)
- Other Files

The main content area shows the 'paper.html' file, which contains the title 'A Standard for the Scholarly Citation of Archaeological Data' by Ben Marwick and Suzanne E. Pilaar Birch, dated 17 April, 2018. The abstract discusses the challenges of data sharing in archaeology and the need for a standard. The bottom of the page notes that data are the building blocks of research and that this paper aims to present a standard for the scholarly citation of archaeological data.

The right sidebar shows the 'Re-Run' section with a timeline of events:

- Apr 18, 2018: Published Version 1.0 (Currently viewing)
- Ben Marwick committed Apr 18, 2018
- Author ran Apr 17, 2018 (0:00:52)
- Sep 28, 2017: Created capsule

At the bottom right, a vertical label 'Reproducibility' is visible.

Publishing reproducible R notebooks

- <https://codeocean.com/capsule/5777882/tree/v1> for .Rmd
- <https://codeocean.com/capsule/0129473/tree> for *.R.

The screenshot displays the Code Ocean web interface for a reproducible R notebook. The top navigation bar includes the Code Ocean logo, a 'Shared' status, the project name 'tempcon_pub_R' by 'Halle R. Dimsdale-Zucker et al.', and buttons for 'Capsule', 'File', 'Edit', 'View', 'Tabs', 'Settings', and 'Help'. A 'Start editing' button is also present.

The left sidebar shows a file explorer with 'Core Files' and 'Other Files'. The 'Core Files' section includes 'environment' (3.05 KB), 'code' (1.75 MB), 'matlabfunctions' (1.31 MB), 'mri_analyses' (8.18 KB), 'tempcon' (441.33 KB), 'config.yml' (403 B), 'LICENSE' (1.05 KB), and 'run.sh' (3.03 KB). The 'Other Files' section includes '.Rproj.user' (1.93 MB) and 'capsule.Rproj' (205 B).

The main area shows the 'run.sh' script, which is a bash script that sets up the environment and runs R scripts. The script includes comments and commands for setting the shell, running R scripts, and generating output files. The script is as follows:

```
1 #!/bin/bash
2 set -ex
3
4 # --- R analyses (behavior and prepping MRI data for modeling) ---
5 # R scripts like to be run from their code directory so cd then run scripts
6 cd ../tempcon
7
8 # this script will load in the presentation-format behavioral data and put it into a nice(r) R
9 # format to work with
10 Rscript -e "rmarkdown::render(input = 'load_data.R', \
11   output_dir = '../results/')"
12
13 # now, let's "tidy" up the behavioral data
14 Rscript -e "rmarkdown::render(input = 'behav-tidy.R', \
15   output_dir = '../results/')"
16
17 # create onset and trial information files (these are used in the fMRI analyses)
18 Rscript -e "rmarkdown::render(input = 'create-onset-files.R', \
19   output_dir = '../results/')"
20
21 # run some stats on the behavior and generate plots
22 Rscript -e "rmarkdown::render(input = 'behav-stats.R', \
23   output_dir = '../results/')"
24
25 # --- matlab analyses (MRI-data related) ---
26 # change directories so matlab is happy
27 cd ../mri_analyses
28
29 # load matlab paths and generate single trial regressors
30 matlab -nodisplay -nosplash -nosoftawareopen -r "addpath(genpath('../code/matlabfunctions'));
31   RSA_generate_single_trial_regressors"
32
33 # next steps aren't feasible to represent here, but they include (scripts for these steps are
34 # included in the GitHub repo: https://github.com/hallez/tempcon_pub):
35 # 1. single trial modeling ('RSA_single_trial_models_batch.n')
36 # 2. ROI-specific things (e.g., extracting from tracings, getting ROIs into the correct space,
37 # etc.)
38 # 3. extracting values w/in the ROIs of interest from the single trial betas
39 ('RSA_extract_betas_from_ROIs.m')
40
41 # --- switch back to R for the meat of the RSA ---
42 cd ../tempcon
43
44 # generate the masks for grabbing trials of interest
```

The right sidebar shows the 'Reproducible Run' section, which includes a timeline of runs. The first run is 'Published Version 1.0' on May 6, 2019, with a button to 'Go To Published Capsule'. The second run is 'Halle Dimsdale-Zucker ran' on May 5, 2019, at 1:25:35. This run includes a list of files generated, such as 'behav-stats.html' (1.49 MB), 'behav-tidy.html' (1.28 MB), 'create-onset-files.html' (950.9 KB), 'load_data.html' (1.23 MB), 'mixed-models-btwn-r...' (2.25 MB), 'output' (67.56 KB), 'perms_list-samediff-h...' (1.52 MB), 'rsa-generate-masks...' (1.29 MB), and 'rsa-tidy-data-btwn-runs...' (2 MB). The third run is 'Halle Dimsdale-Zucker committed' on May 5, 2019. The fourth run is 'Halle Dimsdale-Zucker ran' on May 5, 2019, at 0:06:23, with a button to 'Cloud Workstation output'. The fifth run is 'Halle Dimsdale-Zucker' on May 5, 2019.

Publishing reproducible Jupyter Notebooks

- Notebook + environment + nbconvert = a rendered HTML
- <https://codeocean.com/capsule/6314882/tree/v1>

← → ↺ <https://codeocean.com/capsule/6314882/tree/v1> 🔍 ☆ 🌐 📄 👤 ⋮

CO Published Identifying Gene Expression Programs of Cell-type Identity and Cellular Activity with Sin... (Dylan Kotliar et al.)

Capsule File Edit View Tabs Settings Help

Files

- Core Files
- environment 2.01 KB
- code 21.32 MB
 - analysis 21.23 MB
 - code 79.26 KB
 - Download_Pregenerated_Int... 3.55 KB
 - install_utils.sh 314 B
 - LICENSE 1.04 KB
 - matplotlibrc 213 B
 - README.md 5.95 KB
 - run.sh 1.16 KB
 - Table_Of_Contents.ipynb 4.03 KB
- data Manage Datasets
 - Part1_Simulations 14.15 GB
 - Part2_Organoids 2.02 GB
 - Part3_VisualCortex 2.92 GB
 - LICENSE 6.4 KB
 - .gitignore 7 B
- Other Files

Commands

Tabs

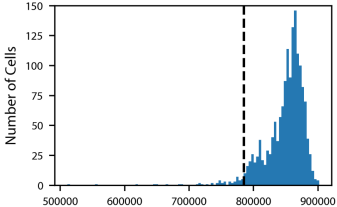
README.md x Step1_Preproc x Step1_Simulat x Step2_Preproc x

in the previous filters

```
In [16]: TPM = pd.read_csv('../.../data/Part3_VisualCortex/GSE71585_RefSeq_TPM.csv', index_col=0).T
```

```
In [17]: (fig,ax) = plt.subplots(1,1, figsize=(3,2), dpi=400)
remaining_tpm_cutoff = 785000
remaining_TPM = TPM.loc[counts_per_kb.index, counts_per_kb.columns].sum(ax
is=1)
_ = ax.hist(remaining_TPM, bins=100)
ax.vlines(x=remaining_tpm_cutoff, ymin=0, ymax=150, linestyle='--')
ax.set_ylim([0,150])
ax.set_xlabel('Remaining TPM After Filters')
ax.set_ylabel('Number of Cells')
```

```
Out[17]: Text(0,0.5,'Number of Cells')
```



Re-Run

Timeline

Nov 20, 2018 Published Version 1.0 Currently viewing

Dylan Kotliar committed Nov 20, 2018

Version 1.0

Author ran Nov 19, 2018 12:33:07

Published Result

- figures
 - Output 20.42 KB
 - Step0_Estimate_Si... 253.93 KB
 - Step1_Preprocess... 384.92 KB
 - Step1_Preprocess... 419.64 KB
 - Step1_Simulate.ht... 309.13 KB
 - Step2_Preprocess... 555.47 KB
 - Step2_Preprocess... 308.63 KB
 - Step2_Run_cNMF_Q... 9.29 MB
 - Step3_Run_cNMF_H... 1.07 MB
 - Step3a_Run_cNMF... 362.09 KB
 - Step3b_Run_dCA.h... 323.1 KB
 - Step3c_Run_cLDA... 324.23 KB
 - Step3d_Run_PCA... 256.91 KB
 - Step3e_Run_Clust... 273.09 KB
 - Step3f_RefSeq_Spect... 258.71 KB




Interactive sessions

- <https://codeocean.com/capsule/8962292/tree/v2>
- Jupyter
- JupyterLab
- Time for a live demonstration

Questions?

- How is this different than Binder?
- What is the uploading process like?
- How are dependencies managed?
- Is this exportable?
- Whom is Code Ocean for? (All scientists)
- What are the most pressing issues in computational reproducibility? (e.g., big data, confidential data, inferring system level dependencies from listed scientific libraries, adjudicating between many competing worthy aims, what language should we all be using. . .) ## Reference Slide 1: Publishing on Code Ocean:
- <https://codeocean.com/2018/10/16/the-contact-hypothesis-re-evaluated-colon-code-and-data/code>
- Will have a DOI and link to your article's metadata

← → ↺ <https://codeocean.com/2018/10/16/the-contact-hypothesis-re-evaluated-colon-code-and-data/metadata> 🔍 ☆ 🌐 📄 👤 ⋮

 v5 > The contact hypothesis re-evaluated: code and data (Betsy Levy Paluck et al.) [Switch to Old Editor](#) [Back to Capsule](#)  

Basic Info

Language

Stata

Compute Capsule DOI  <https://doi.org/10.24433/CO.f152260c-bebb-4157-a640-44579452b4e4.v5>

Reference Slide 2: Embedding on webpages & within articles

- You can also embed your published capsule in your article's HTML page or on your personal webpage, a la <https://ieeexplore.ieee.org/document/8410389/algorithms#algorithms>:

[//ieeexplore.ieee.org/document/8410389/algorithms#algorithms](https://ieeexplore.ieee.org/document/8410389/algorithms#algorithms):

explore.ieee.org/document/8410389 67% ... ☆ 🔍 See

Keywords

Metrics

Code & Datasets

Code

Dataset

This article contains code hosted on IEEE's partner, Code Ocean, a cloud-based computational reproducibility platform that enables users to run, modify, and download code from IEEE Xplore articles. A Code Ocean user account is required to run and modify code within the widget below.

Code: On Writing Reproducible and Interactive Papers Python

The screenshot displays the Code Ocean interface for a capsule titled "On Writing Reproducible and Interactive Papers" by Mandar Chitre. The interface is divided into several sections: a top navigation bar with "Capsule", "File", "Edit", "View", "Tabs", "Settings", and "Help"; a left sidebar with "Files", "Commands", and "Tabs"; a main content area showing the "Editorial" text; a right sidebar with "Reproducibility" and a list of files including "output" (27.61 KB), "editorial.pdf" (123.91 KB), and "editorial.tex" (27.05 KB). A "Run" button is prominently displayed, and the capsule is marked as "Run 5395314" with a timestamp of "0:00:11". The capsule is verified as published by Mandar Chitre on June 13, 2018.