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Motivation: Prevent Loss of Knowledge

Every semester, graduate students around **the world** take an Empirical/Applied [...] Economics course. A typical assignment consists of reproducing the results of a paper and, possibly, testing the robustness of its results.

Stage	New Knowledge
Scope (select and verify)	Data and code exist?
Assess	Degree of reproducibility for specific part of the paper
Improve	E.g. fixed paths, libraries, added missing files, etc.
Test robustness	Results are robust to additional specifications

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Context for ACRE

- American Economics Association (AEA) creates first data policy in 2006.
 - Must publish some data (waivers available)
- AEA updates policy in 2019.
 - Must post all data and code. Publication is conditional on verifying reproducibility (if confidential: must document extensively)
 - A new requirement is to post all cleaning code, even for data that is not public
 - See the AEA Data Editor Website for more information
- We should expect high levels of computational reproducibility after 2019 (AEA Journals).
- We should not demand 100% reproducibility before, but we could identify the gaps and try to improve some.

Beyond Binary Judgments

Reproductions can easily gravitate towards adversarial exchanges.

- Early career researcher (ECR) have incentives to emphasize unsuccessful reproductions
- Original authors have a more senior position and can use it to deter in-depth reproductions from ECRs.
- The media also focuses on eye-catching headlines

Our approach:

We do not want to say

"Paper X is (ir)reproducible"

We do want to say

"Paper X's result Y has a high/low **level** of reproducibility according to **several** reproduction attempts. Moreover, **improvements** have been made to the original reproduction package, **increasing** its reproducibility to a higher level"

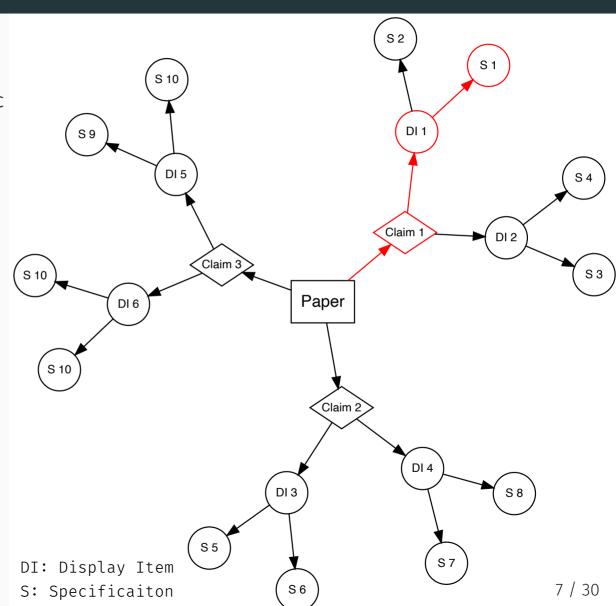
Our Framework

Each **reproduction attempt** is centered around scientific **claims**

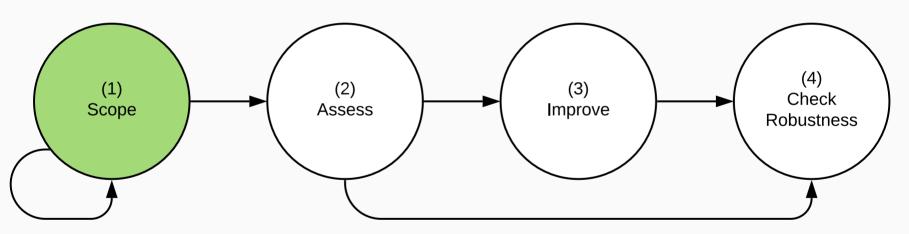
One paper can contain several claims.

Each claim may be supported by various **display items**: tables, figures & inline results.

A reproduction attempt is at the claim level, and reproducers must record their **specifications** of interest.



Stages



Scoping

- 1. Select or be assigned a candidate paper
- 2. Check ACRE Platform for previous entries and verify availability of reproduction package (RP)
- 3. If no RP, leave a short record, and repeat with a different candidate paper
- 4. Once RP is found then candidate becomes declared paper
- 5. Only then: read the paper and select claim(s), display items and specification to reproduce

Box 1: Summary Report Card for

ACRE Paper Entry

Title: Sample Title

Authors: Jane Doe & John Doe

Original Reproduction Package

Available: URL/No

[If "Yes"]

Additional Reproduction Packages:

Number (eg., 2)

[If "No"]

Contacted Authors?: Yes/No

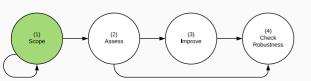
[If "Yes(contacted)"]

Type of Response: Categories (6).

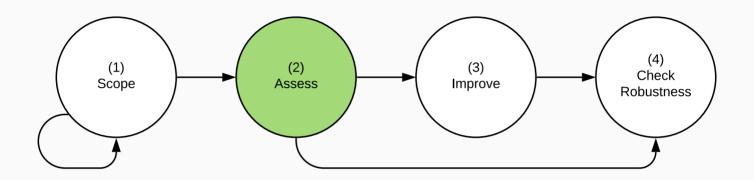
Authors Available for Further

Questions for ACRE Reproductions:

Yes/No/Unknown



Assessment



Two main parts for assessment:

- 1. Find all the elements behind a display item
- 2. Score the reproducibility of that display item

Identify All the Elements Behind a Display Item

Reproducers will be asked to draw a clear connection to the raw data sources mentioned in the paper and the display item under reproduction.

Data sources

Connect the data sources in the paper's text with specific raw data files.

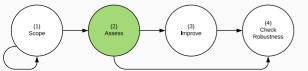
Analytic data sets

Describe each analytic data file.

Code files

Inspect all code files and record all their inputs and outputs.

With all the information recorded above, reproducers can use the ACRE Diagram Builder to generate a **reproduction tree**.

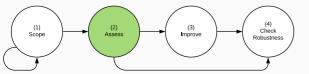


Reproduction Tree

```
table1.tex
   [code] analysis.R
       | analysis data.dta
          [code] final merge.do
              cleaned 1 2.dta
                [code] clean merged 1 2.do
                    merged 1 2.dta
                        [code] merge 1 2.do
                            | cleaned 1.dta
                             [code] clean_raw_1.py
                                  raw 1.dta
                            | cleaned 2.dta
                               [code] clean raw 2.py
                                  ___raw_2.dta
              cleaned 3 4.dta
                 [code] clean merged 3 4.do
                    merged 3 4.dta
                        [code] merge 3 4.do
                            | cleaned 3.dta
                             |___[code] clean_raw_3.py
                                  raw 3.dta
                            | cleaned 4.dta
                               [code] clean_raw_4.py
                                  raw 4.dta
```

Levels

```
Levels of Computational Reproducibility
                (P denotes "partial", C denotes "complete")
                          Availability of materials, and reproducibility
                          |Analysis| Analysis| | Cleaning| Raw
                          L1: No materials.....
L2: Only code ..... ✓
L3: Partial analysis data & code.
L4: All analysis data & code..... ✓
L5: Reproducible from analysis ... | ✓
L6: Some cleaning code..... ✓
L7: All cleaning code..... ✓
L8: Some raw data..... ✓
L10:Reproducible from raw data... |
```

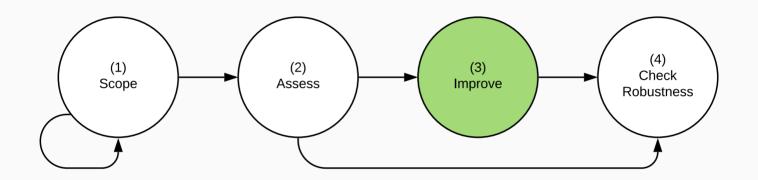


Levels: Proprietary/Confidential Data

```
Levels of Computational Reproducibility
                      with Proprietary/Confidential Data
                   (P denotes "partial", C denotes "complete")
                               Availability of materials, and reproducibility
                                      | Instr. | | Instr. |
                              |Analysis| Analysis| | Cleaning| Raw
                                      | Data | CRA | Code
L1: No materials.....
L2: Only code ..... ✓
L3*: Partial analysis data & code ✓
L4*: All analysis data & code.... ✓
L5*: Proof of third party CRA.... ✓
L6: Some cleaning code..... ✓
L7: All cleaning code..... ✓
L8*: Some instr. for raw data.... ✓
L9*: All instr. for raw data..... ✓
L10*: Proof of third party CRR.... ✓
```



Improvements

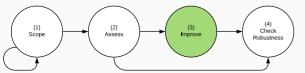


Three types of improvements:

- 1. Improvements at the paper level
- 2. Improvements at the display-item level
- 3. Specific future improvements

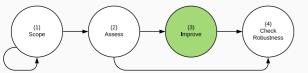
Improvements: Paper-level

- Use version control software (Git/Github).
- Improve documentation: comments, indentations, object names, etc.
- Re-organize the reproduction package into a set of folders and sub-folders that follow standardized best practices, and add a master script that executes all the code in order, with no further modifications. See AEA's reproduction template.
- Literate programming environment (e.g., Jupyter notebooks, RMarkdown)
- Re-write code using a differenet statistical software (ideally open source, like R, Python, or Julia).
- Set up a computing capsule (e.g., Binder and Code Ocean).

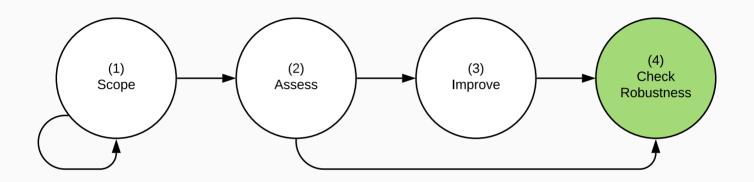


Improvements: Display item-level

- Adding missing raw data: files or meta-data
 - Example: "Add raw temperature and relative humidity data"
- Adding missing analytic data files
 - Example: "Copy the row files from Data folder into new Analysis\trade cost\Input"
- Adding missing analysis or cleaning code
 - Example: "Replaced broken Wald bootstrap code with updated code/command"
- Debugging code
 - Example: "was counting each group 4 times in round 1, so fixed that"
- Propose specific future improvements



Robustness Checks



Two main parts for robustness:

- 1. Increase the number of possible robustness checks
- 2. Justify the appropriateness of a specific test

Robustness

Robustness checks: any possible change in a computational choice, both in data analysis and data cleaning

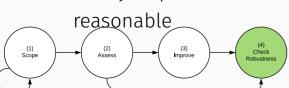
Reasonable specifications (Simonsohn et.

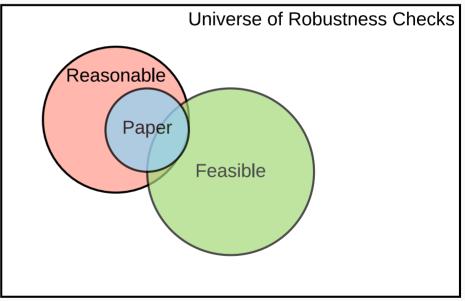
al., 2018):

- 1. Sensible tests of the research question
- 2. Expected to be statistically valid, and
- 3. Not redundant with other specifications in the set.

Reproducers will be able to record two types of contributions:

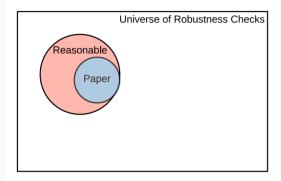
- Mapping the universe of robustness checks
- Justify a specific robustness check as



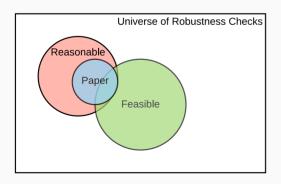


Robustness & Reproducibility

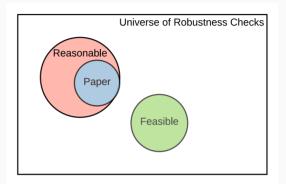
Robustness with level 1



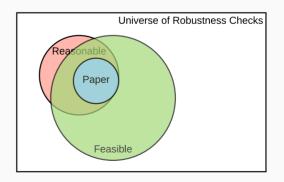
Robustness with levels 5-9

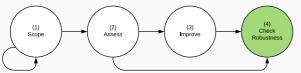


Robustness with levels 2-4



Robustness with level 10





Promoting a Constructive Exchange

- 1 Contacting the original author(s) when there is no reproduction package
- 2 Contacting the original author(s) to request specific missing items of a reproduction package
- 3 Asking for additional guidance when some materials have been shared
- 4 Response when the original author has refused to share due to undisclosed reasons
- 5 Response when the original author has refused to share due to legal or ethical restrictions of the data
- 6 Contacting the original author to share the results of your reproduction exercise
- 7 Responding to hostile responses from original authors

Under development: sample responses form authors to reproducers

Example 1: There is no reproduction package

```
Subject: Reproduction package for ["Title of the paper"]
Dear Dr. [Lastname of Corresponding Author],
I am contacting you to request a reproduction package for your paper titled [Title] which was published in
[Reference]. A reproduction package may contain (raw and/or analytic) data, code, and other documentation that
makes it possible to reproduce paper. Would you be able to share any of these items?
I am a [position] at [Institution], and I would like to reproduce the results, tables, and other figures using the
reproduction materials mentioned above. I have chosen this paper because [add context ...]. Unfortunately, I
was not able to locate any of these materials on the journal website, Dataverse [or other data and code
repositories], or in your website.
I will record the result of my reproduction attempt on ACRE [...]. With your permission, I will also record the
materials you share with me, which would allow access for other reproducers and avoid repeated requests
directed to you. Please let me know if there are any legal or ethical restrictions that apply to all or parts of the
reproduction materials so that I can take that into consideration during this exercise.
In addition to your response above, would you be available to respond to future (non-repetitive) inquiries from me
or other reproducers conducting an ACRE excercise? Though your cooperation with my and/or any future request
would be extremely helpful, please note that you are not required to comply.
Since I am required to complete this project by [date], I would appreciate your response by [deadline].
Let me know if you have any questions. Please also feel free to contact my supervisor/instructor [Name (email)] for
further details on this exercise. Thank you in advance for your help!
Best regards,
```

[Reproducer]

Example 1: Following up on additional materials

Template email:

Subject: Clarification for reproduction materials for ["Title of the paper"]

Dear Dr. [Lastname of Corresponding Author],

Thank you for sharing the materials. They have been immensely helpful for my work.

Unfortunately, I ran into a few issues as I delved into the reproduction exercise, and I think your guidance would be helpful in resolving them. [Describe the issues and how you have tried to resolve them. Describe whatever files or parts of the data or code are missing. Refer to examples 1 and 2 below for more details].

Thank you in advance for your help.

Best regards,

[Reproducer]

An example of well described issues:

Specifically, I am attempting to reproduce OUTPUT X (e.g., table 1, figure 3). I found that the following components are required to reproduce to reproduce OUTPUT X:

```
OUTPUT X

——[code] formatting_table1.R

——output1_part1.txt

| ——[code] output_table1.do

| ——[data] analysis_data01.csv

| ——[code] data_cleaning01.R*

| ——[data] UNKNOWN

——output1_part2.txt

——[code] output_table2.do

——[code] output_table2.csv

——[data] analysis_data02.csv

——[code] data_cleaning02.R

——[data] admin_01raw.csv*
```

I have marked with an asterisk (*) the items that I could not find in the reproduction materials: **data_cleaning01.R** and **admin_01raw.csv**. After accessing these files, I will also be able to identify the name of the raw data set required to obtain output1_part1.txt. This is to let you know that I may need to contact you again if I cannot find this file (labeled as **UNKNOWN** above) in the reproduction materials.

I understand that this request will require some work for you or somebody in your research group, but I want to assure you that I will add these missing files to the reproduction package for your paper on the ACRE platform. **Doing this will ensure that you will not be asked twice for the same missing file.**

Ok, I get it. But what is in for me?

- Standardized homework/project: everything is set up in terms of structure and deliverables.
- Easy to grade (homework format).
- Easy to guide and oversee (undergraduate dissertation format).
- Easy to setup as an independent study.
- Reduces duplication of requests to authors.
- Facilitates a constructive exchange of ideas.
 - When emailing authors.
 - When discussion reproduction attempts.
- Personal satisfaction that you're contributing a public good to the profession!

Easy to grade: report 1

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Easy to grade: report 1

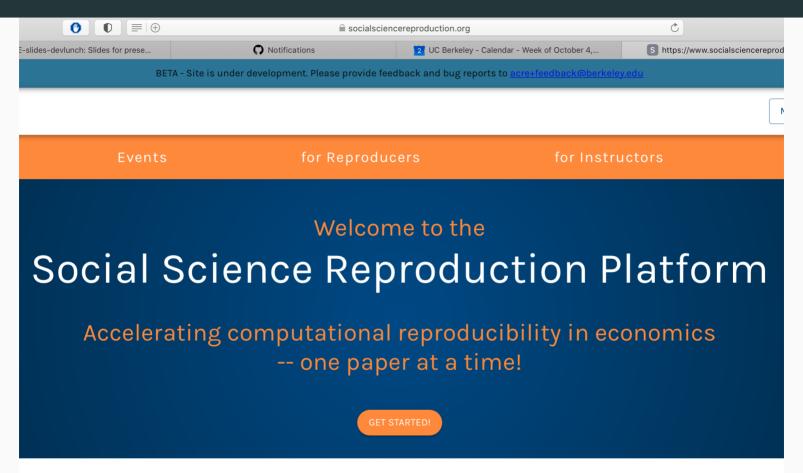
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socialsciencereproduction.org



Purpose

On the Social Science Reproduction Platform, you can record and review **verifications and improvements** to the **computational reproducibility** of published social science work.

This open source platform was developed by the Berkeley Initiative for Transparency in the Social Sciences (BITSS) in collaboration with the American Economic Association Data Editor.

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