

Automated, semi-automated, and manual extraction of numerical data from scientific images with the *juicr* package for *R* [BETA]

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University of South Florida, February 18th 2026 (vignette v. 0.2 for *juicr* v. 0.2)

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Introduction

The **juicr** package for *R* contains tools for facilitating the extractions of numerical data from scientific images – like scatter-plots, bar-plots, and other charts/figures found in publications. Below is a description of functionalities and layout.

Updates to this vignette will be posted on our [research webpage at USF](#).

For the source code of **juicr** see: <https://CRAN.R-project.org/package=juicr>.

Acknowledgements

I thank everyone who watched my *YouTube* course [Hard-boiled Synthesis](#) and reached out to me about using **juicr** – you gave me the final push to complete this old project I started way back and abandoned in 2017!

How to cite? TBA, but for this beta version maybe:

Lajeunesse, M.J. (2021) Squeezing data from scientific images with the **juicr** package for *R*. *R* package, v. 0.1. [CRAN](#)

Installation and Dependencies

juicr has an external dependency that needs to be installed and loaded prior to use in *R*. This is the **EBImage** *R* package (Pau *et al.* 2010) available only from [Bioconductor](#) repository.

To properly install **juicr**, use the following script in *R*:

```
# first Load Bioconductor resources needed to install the EBImage package
# and accept/download all of its dependencies
install.packages("BiocManager");
BiocManager::install("EBImage")

# then Load juicr
library(juicr)
```

Finally for Mac OS users, installation is sometimes not straightforward, as the `GUI_juicr()` requires the Tcl/Tk GUI toolkit to be installed. You can get this toolkit by making sure that the latest X11 application (xQuartz) is installed from here: <https://www.xquartz.org/>.

Report a bug? Have comments or suggestions?

Please email me any bugs, comments, or suggestions and I'll try to include them in future releases: lajeunesse@usf.edu. Also try to include **juicr** in the subject heading of your email. Finally, I'm open to almost anything, but expect a lag before I respond and/or new additions are added.



GUI layout and loading images

When running **juicr** without a file specified, the layout is simple:

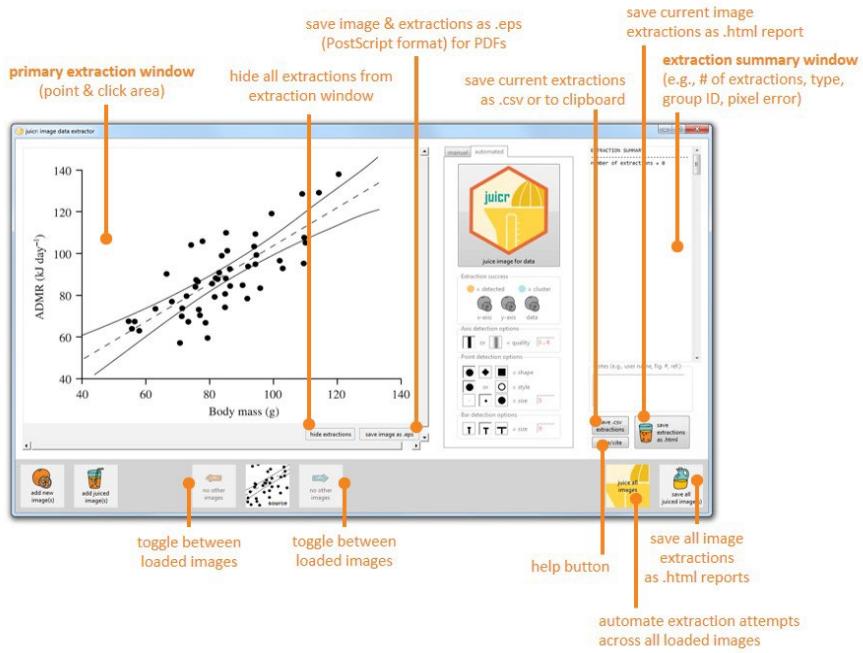


select single or multiple image files
select single or multiple pre-extracted **juicr** files (format: *_juicr.html)

Images can be loaded into **juicr** using the *add new image(s)* button. Alternatively one or many images (as a vector of file name strings) can be included via console:

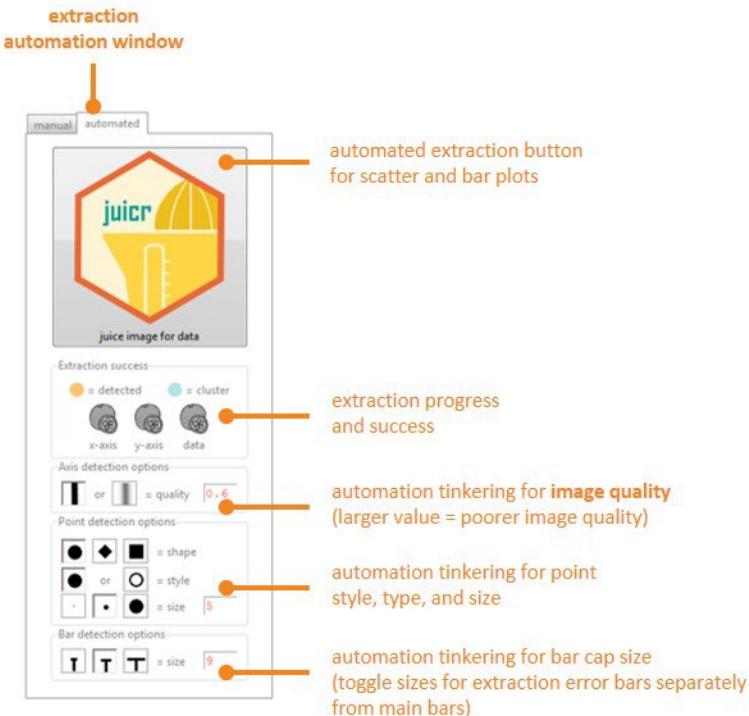
```
# then load juicr
library(juicr)
GUI_juicr("Kam_et_al_2003_Fig2.jpg")
# or many files
GUI_juicr(c("Kam_et_al_2003_Fig2.jpg", "Kortum_and_Acymyan_2013_Fig4.jpg"))
```

If an image is loaded, the main window will look like this:



Automated extraction functionality

The center section of **juicr** includes two options: *automated* or *manual* tools for extracting data. The automated tools include:

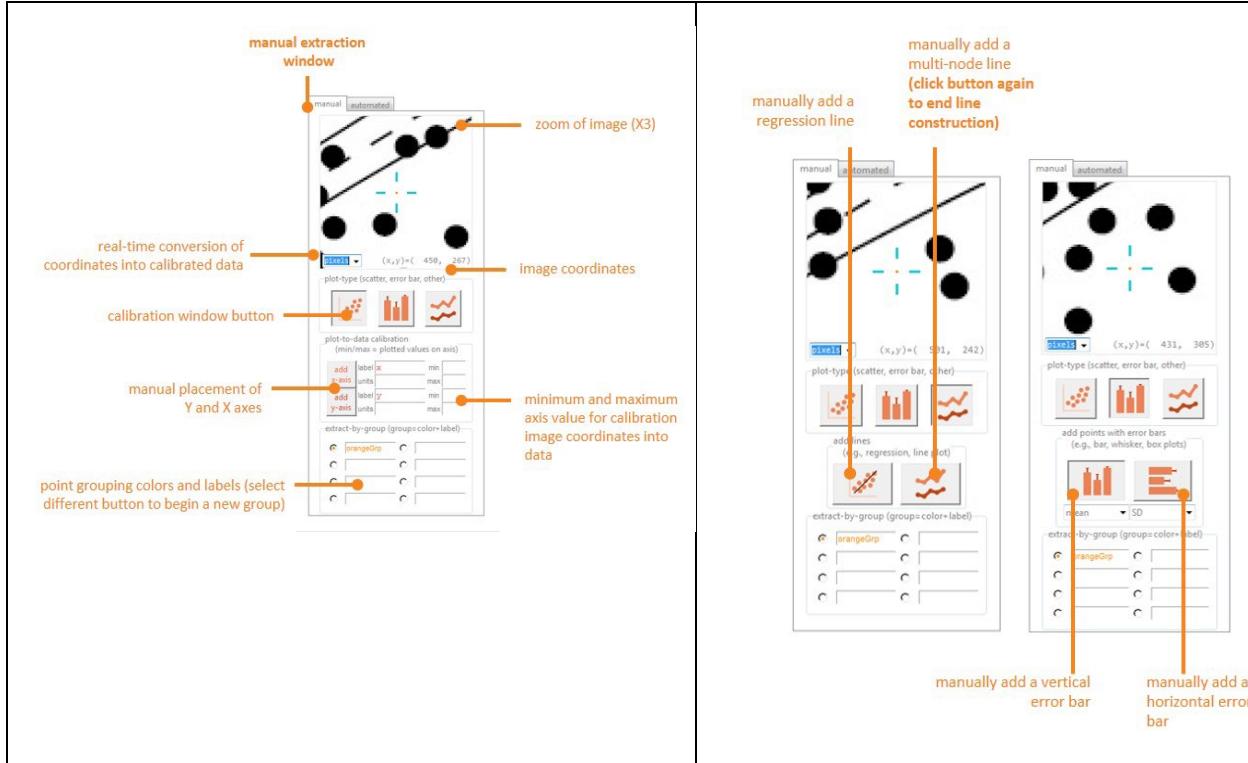


By pressing the large **juicr-hex** button, it will first determine automatically plot type (e.g., scatter or bar for now), and then begin extracting data from the image. Successes of extractions, such as whether the y-axis was detected, will be presented as an orange (detected), and failures as gray (not-detected). Also included a suite of semi-automated tinkering options to help **juicr** detect the desired image objects.



Manual extraction functionality

A large diversity of manual extraction tools are available:



Saving extractions and generating reports

Currently, **juicr** offers extractions to be saved as .csv files, copied into clipboards, saved as postscript .eps files, or as a fully-embedded and standalone .html file that retains all information of extractions, **juicr** setup, and image modifications for permanent and replicable storage of data.

Here is an example of a report:

http://lajeunesse.myweb.usf.edu/juicr/Kam_et_al_2003_Fig2_juicr.html