- 1 Reproducible, flexible and high-throughput data extraction from primary
- 2 literature: The metaDigitise R package
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8 Abstract

- 1. Research synthesis, such as comparative and meta-analyses, requires the
 extraction of effect sizes from primary literature, which are commonly calculated
 from descriptive statistics. However, the exact values of such statistics are
 commonly hidden in figures.
 - 2. Extracting descriptive statistics from figures can be a slow process that is not easily reproducible. Additionally, current software lacks an ability to incorporate important meta-data (e.g., sample sizes, treatment / variable names) about experiments and is not integrated with other software to streamline analysis pipelines.
- 3. Here we present the R package **metaDigitise** which extracts descriptive statistics such as means, standard deviations and correlations from four plot types: 1) mean/error plots (e.g. bar graphs with standard errors), 2) box plots, 3) scatter plots and 4) histograms. **metaDigitise** is user-friendly and easy to learn as it interactively guides the user through the data extraction process. Notably, it enables large-scale extraction by automatically loading image files, letting the user stop processing, edit and add to the resulting data-frame at any point.
 - 4. Digitised data can be easily re-plotted and checked, facilitating reproducible data extraction from plots with little inter-observer bias. We hope that by making the process of figure extraction more flexible and easy to conduct it will improve the transparency and quality of meta-analyses in the future.
- **Keywords:** meta-analysis, comparative analysis, data extraction, R, reproducibility,
- 30 figures, images, descriptive statistics

31 1 Introduction

In many different contexts, researchers make use of data presented in primary 32 literature. In the fields of ecology and evolution (E&E), these data are most commonly 33 used for comparative and meta-analyses. The use of meta-analysis in E&E in 34 particular, is rapidly growing, not only in terms of the number of meta-analyses (in 35 plant ecology alone the yearly number of published meta-analyses doubled from 2006 to 36 2012 (20-40); Koricheva & Gurevitch (2014)), but also in terms of their size (a recent 37 38 meta-analysis, for example, included 6440 effect sizes from 175 publications; Noble, Stenhouse & Schwarz (2018)). Meta-analyses are extremely important in providing a 39 40 means of quantitatively synthesizing experimental and/or observational studies to evaluate empirical support for fundamental theory in E&E (Gurevitch et al., 2018). 41These techniques rely heavily on descriptive statistics (e.g. means, standard deviations 42(SD), sample sizes, correlation coefficients) extracted from primary literature. As well 43 as being presented in the text or tables of research papers, descriptive statistics are 44 frequently presented in figures. For example, in some cases up to 42% of the papers 45 used have one or more figures for data extraction (Noble, Stenhouse & Schwanz, 2018). 46 These data often need to be manually extracted using digitising programs. 47 48 Although there are several tools that extract data from figures, including both standalone programs and R packages (reviewed in Table 1), these tools do not cater to 49 the general needs of meta-analysts for four main reasons (here we focus on 50 meta-analysis, although many points apply to extraction for comparative analysis). 51 First, although meta-analysis is an important tool in consolidating the data from 52 multiple studies, many of the processes involved in data extraction are opaque and 53 difficult to reproduce, making extending or replicating studies problematic. Having a 54 tool that facilitates reproducibility in meta-analyses will increase transparency and aid 55 in resolving the reproducibility crises seen in many fields (Peng, Dominici & Zeger, 56 2006; Peng, 2011; Parker et al., 2016). Second, digitising programs do not allow the 57

integration of metadata at the time of data extraction, such as experimental group or 58 variable names, and sample sizes. This makes the downstream calculations laborious, as 59 information has to be added later, typically using different software. Third, existing 60 programs do not import sets of images for the user to systematically work through. 61 Instead they require the user to manually import images and export the resulting 62 digitised data into individual files one-by-one. These data often subsequently need to be 63 imported and edited using different software. Finally, digitising programs typically only 64 65 provide the user with calibrated x,y coordinates from imported figures, and do not differentiate between common plot types that are used to present data. Consequently, a 66 large amount of additional data manipulation is required, that is different across plots 67 types. For example, in E&E data are commonly presented in plots with means and 68 standard errors or confidence intervals (Figure 1A), from which the user wants a mean 69 70 and SD for each group presented. From x,y coordinates, users must manually discern between mean and error coordinates and assign points to groups. The error then needs 71 to be calculated as the deviation from the mean, and then transformed to SD, according to the type of error presented. Histograms and box plots are also frequently used in 73E&E to presented data, and whilst their downstream calculations are even more 74laborious, there are few (if any; see Table 1) tools to extract data from these plot 75types. 76 Data extraction from figures is therefore a time-consuming process as existing software 77 does not provide an optimized, reproducible research pipeline to facilitate data 78 extraction and editing. Given the ubiquity of the R platform in E&E, and that it hosts 79 the most popular meta-analysis software in E&E (e.g., metafor (Viechtbauer, 2010) and 80 MCMCglmm (Hadfield, 2010)), it is highly likely to be used for some (if not all) stages 81 of the research synthesis process. It is therefore important to have comprehensive, 82 robust and flexible digitisation capabilities in R to make the process of figure extraction 83 more streamline, transparent and easier to reproduce. Here, we present an interactive R 84 package, **metaDigitise** (available on CRAN), which is designed for large scale, 85

reproducible data extraction from figures, specifically catering to the the needs of meta-analysts. To this end, we provide tools to extract data from common plot types in 87 E&E (mean/error plots, box plots, scatter plots and histograms, see Figure 1). 88 metaDigitise operates within the R environment making data extraction, analysis and 89 export more streamlined. The necessary calculations are carried out on calibrated data 90 immediately after extraction so that comparable descriptive statistics can be obtained 91 quickly. Summary data from multiple figures is returned into a single data frame which 92 93 can be can easily exported or used in downstream analysis within R. Completed digitisations are automatically saved for each figure, meaning users can redraw their 94 digitisations (along with metadata) on figures, make corrections and access calibration 95 and processed (i.e., summarised) data. This makes sharing figure digitisation and 96 97 reproducing the work of others simple and easy, and allows meta-analyses to be 98 updated more efficiently.

99 2 metaDigitise and Reproducibility

The metaDigitise package has one main function, metaDigitise(), which interactively 100 takes the user through the process of extracting data from figures (see Supplementary 101 102 Material S1 for a full tutorial). Running metaDigitise() presents the user with three options; 'Process new images', 'Import existing data' or 'Edit existing data', which can 103 104 be used during and after digitisation to execute a range of functions (see Figure 1 – 105 'Processing images' is discussed in Section 3, and 'Editing' and 'Importing' in Section 4). metaDigitise() works on a directory containing images of figures copied from 106 primary literature, in .png, .jpg, .tiff, .pdf format, specified to metaDigitise() through 107 108 the dir argument. metaDigitise() recognizes all the images in the given directory and 109 automatically imports them one-by-one, allowing the user to extract the relevant information about a figure as they go. Figures can be organised in different ways for a 110 project, but we would recommend having all figures for one project in a single directory 111

- 112 with an informative and unambiguous naming scheme (e.g. paper_figure_trait.png).
- 113 This expedites digitisation by preventing users from having to constantly change
- 114 directories and / or open new images.
- 115 The data from each completed image is automatically saved as a metaDigitise object
- in a separate .RDS file to a caldat folder that is created within the parent directory
- 117 when first executing metaDigitise(). These files enable re-plotting and editing of
- 118 images at a later point (see below). When run, metaDigitise() also identifies the
- images within a directory that have been previously digitised and only imports new
- 120 images to process. The data of all images is then automatically integrated into the final
- 121 output. This means that all figures do not need to be extracted at one time and new
- 122 figures can be added to the directory as the project develops.
- 123 The complete digitisation process can be reproduced at a later stage, shared with
- 124 collaborators and presented as supplementary materials for a publication, regardless of
- 125 the computer it is run on. To update an analysis, new figures can simply be added to
- 126 the directory and metaDigitise() run to incorporate the new data.

127 3 Image Processing

- 128 Selecting 'Process New Images', after running metaDigitise(), starts the digitisation
- 129 process on images within the directory that have not previously been digitised. For all
- 130 plot types, metaDigitise() requires the user to calibrate the axes in the figure, by
- 131 clicking on two known points on the axis in question, and entering the value of those
- points (Figure 1). metaDigitise() then calculates the value of any clicked points in
- 133 terms of the figure axes. This is based on the calibration used in the **digitize** R package
- 134 (Poisot, 2011). For mean/error and box plots, only the y-axis is calibrated (Figure 1),
- 135 assuming the x-axis is redundant. For scatter plots and histograms both axes are
- 136 calibrated (Figure 1).

Calibration of points in figures from older, scanned publications can be problematic, as 137 the figures may not be perfectly orientated. metaDigitise() allows users to rotate the 138 image (Figure S2A,B). Furthermore, mean/error plots, box plots and histograms, may 139 be presented with horizontal bars. metaDigitise() assumes that bars are vertical, but 140 141 allows the user to flip the image to make the bars are vertical (Figure S2C,D). metaDigitise also allows back calculation of data presented on log axes. 142 metaDigitise recognises four main types of plot; Mean/error plots, box plots, scatter 143 plots and histograms (Figure 1). All plot types can be extracted in a single call of 144 145 metaDigitise() and integrated into one output. Alternatively, users can process different plot types separately, using separate directories. All four plot types are 146 extracted slightly differently (outlined below). Upon completing all images, or quitting, 147 either summarised or calibrated data is returned (specified by the user through the 148 149 summary argument). Summarised data consists of a mean, SD and sample size, for each 150 identified group within the plot (should multiple groups exist). In the case of scatter plots, the correlation coefficient between x and y variables within each identified group 151 152 is also returned. Calibrated data consists of a list with slots for each of the four figure types, containing the calibrated points that the user has clicked. This may be 153

155 3.1 Mean/Error and Box Plots

particularly useful in the case of scatter plots.

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metaDigitise() handles mean/error and box plots in a very similar way. For each mean/box, the user enters group name(s) and sample size(s). If the user does not enter a sample size at the time of data extraction (if, for example, the information is not readily available) a SD is not calculated. Sample sizes can, however, be entered at a later time (see next section). For mean/error plots, the user clicks on an error bar followed by the mean. Error bars above or below the mean can be clicked, as sometimes one is clearer than the other. metaDigitise() assumes that the error bars are symmetrical. Points

are displayed where the user has clicked, with the error in a different colour to the mean 163 164 (Figure 1A). The user also enters the type of error used in the figure: SD, standard error (SE) or 95% confidence intervals (CI95). For box plots, the user clicks on the 165 maximum, upper quartile, median, lower quartile and minimum. For both plot types, 166 167 the user can add, edit or remove groups while digitising for when finished. Three functions, error_to_sd(), rqm_to_mean() and rqm_to_sd(), that convert different error 168 169 types to SD, box plot data to mean and box plot data SD, respectively, are also 170 available in the package (see supplements for further details of these conversions).

171 3.2 Scatter plots

Users can extract points from multiple groups from scatter plots. Different groups are 172 plotted in different colours and shapes to enable them to be distinguished, with a legend 173at the bottom of the figure (Figure 1D). Mean, SD and sample size are calculated from 174 175 the clicked points, for each group. Data points may overlap with each other making it impossible to know whether points have been missed. This may result in the sample 176 size of digitised groups conflicting with what is reported in the paper. However, users 177 also have the option to input known sample sizes directly, if required. Nonetheless, it is 178 important to recognise the impact that overlapping points can have on descriptive 179 statistics, and in particular on sampling variance. 180

181 3.3 Histograms

The user clicks on the top corners of each bar, which are drawn in alternating colours (Figure 1C). Bars are numbered to allow the user to edit them. As with scatter plots, if the sample size from the extracted data does not match a known sample size, the user can enter an alternate sample size. The formulas for calculation of mean, SD and sample size are provided in the supplement.

187 4 Importing and Editing Previously Digitised 188 data

metaDigitise is also able to re-import, edit and re-plot previously digitised figures. 189 190 When running metaDigitise(), the user can choose to 'Import existing data', which returns previously digitised data, from a single figure or all figures. Alternately, the 191 getExtracted() function returns the data from previous digitisations, but without user 192 193 interaction, allowing easier integration into larger scripts. 'Edit existing data' allows the user to re-plot or edit information for digitisations that have previously be done. 194 195 Re-plotting digitisations with all metadata is an important reproducibility feature, as it allows users to see exactly what information has been extracted, as well as making it 196 197 easy to spot and data extraction errors.

198 4.1 Adding Sample Sizes to Previous Digitisations

In many cases sample sizes may not be readily available when digitising figures. This information does not need to be added at the time of digitisation. To expedite finding and adding these sample sizes at a later point, metaDigitise() has a specific edit option that allows users to enter previously omitted sample sizes. This first identifies missing sample sizes in the digitised output, re-plots the relevant figures and prompts the user to enter the sample sizes for the relevant groups in the figure.

205 5 Software Validation

In order to evaluate the consistency of digitisation with **metaDigitise** between users, fourteen people digitized sets of 14 identical images created from a simulated dataset (see supplements). We found no evidence for any inter-observer variability in

digitisations for the mean (ICC = 0, 95% CI = 0 to 0.029, p > 0.999), SD (ICC = 0, 209 95% CI = 0 to 0.033, p > 0.999) or correlation coefficient (ICC = 0.053, 95% CI = 0 to 210 211 0.296, p = 0.377). There was little bias between digitised and true values, on average 1.63% (mean = 0.02%, SD = 4.9%, r = -0.03%) and there were small absolute 212 differences between digitised and true values, on average 2.18% (mean = 0.40%, SD = 2135.81%, r = 0.33%) across all three descriptive statistics. SD estimates from digitisations 214 are clearly most error prone. The mean absolute differences for each plot type clearly 215 216 show that this effect is driven by extraction from box plots and histograms (% difference; box plot: 15.81, histogram: 5.21, mean/error: 1.50, scatter plot: 0.43). SD 217218 estimation from box plot descriptive statistics is known to be more error prone, especially at small sample sizes (Wan et al., 2014). 219 220 We also used simulated data to test the accuracy of digitisations with respect to known

values (see supplements). **metaDigitise** was very accurate at matching clicked points to

their true values essentially being perfectly correlated with the true simulated data for

both the x-variable (Pearson's correlation; r > 0.999, t = 2137.4, df = 78, p < 0.001)

and y-variable (r > 0.999, t = 1897.8, df = 78, p < 0.001) in scatterplots.

225 6 Limitations

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226 Although **metaDigitise** is very flexible and provides functionality not seen in any other package, there are some functions that it does not perform (see Table 1). Notably 227228 metaDigitise lacks automated point detection. However, from our experience, manual 229 digitising is more reliable and often equally as fast. Given the variation in image 230 quality, calibration for automatic point detection needs to be done for each figure individually. Additionally, auto-detection often misses points which then need to be 231manually added. Based on tests of **metaDigitise** (see above), figures can be extracted in 232 around 1-2 minutes, including the entry of metadata. As a result, we do not believe 233

that current automated point detection techniques provide substantial benefits in terms 234 of time or accuracy. 235

metaDigitise also (currently) lacks the ability to zoom in on figures. Zooming may 236 enable users to gain greater accuracy when clicking on points. However, from our own 237 experience (see results above), with a reasonably sized screen accuracy is already high, 238 and so relatively little gain is to be had from zooming in on points. 239 240 In contrast to some other packages **metaDigitise** does not extract lines from figures. 241 Although line extraction is not generally necessary in comparative and meta-analysis, outside of these fields researchers may need to extract parameters of a line from a 242figure. Should a user like to extract lines with metaDigitise, we would recommend 243 244 extracting data as a scatter plot, and clicking along the line in question. A model can then be fitted to these points (accessed by choosing to return calibrated rather than 245 summary data) to estimate the parameters needed.

Conclusions 247

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Increasing the reproducibility of figure extraction for meta-analysis and making this 248 laborious process more streamlined, flexible and integrated with existing statistical 249 software will go a long way in facilitating the production of high quality meta-analytic 250 251 studies that can be updated in the future. We believe that **metaDigitise** will improve this research synthesis pipeline, and will hopefully become an integral package that can 252 253 be added to the meta-analysts toolkit.

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263 Author Contributions

- 264 J.L.P. and D.W.A.N. conceived the study and J.L.P., S.N. and D.W.A.N. developed the
- 265 idea. J.L.P. and D.W.A.N. developed the R-package. J.L.P. and D.W.A.N. wrote the
- 266 first draft of the paper and J.L.P., S.N. and D.W.A.N. contributed substantially to
- 267 subsequent revisions of the manuscript and gave final approval for publication.

268 References

- 269 Arizona-Software (2008) GraphClick Software, Version 3.0.
- 270 Bormann, I. (2012) Digitizelt Software, Version 2.0. Braunschweig, Germany.
- 271 Gurevitch, J., Koricheva, J., Nakagawa, S. & Stewart, G. (2018) Meta-analysis and the
- science of research synthesis. *Nature*, **555**, 175–182.
- 273 Hadfield, J.D. (2010) MCMC methods for multi-response generalized linear mixed
- 274 models: The {MCMCglmm} {R} package. Journal of Statistical Software, 33, 1–22.
- 275 Koricheva, J. & Gurevitch, J. (2014) Uses and misuses of meta-analysis in plant
- 276 ecology. *Journal of Ecology*, **102**, 828–844.
- 277 Lajeunesse, M.J. (2016) Facilitating systematic reviews, data extraction, and

- 278 meta-analysis with the metagear package for R. Methods in Ecology and Evolution, 7,
- 279 323–330.
- 280 Noble, D.W., Stenhouse, V. & Schwanz, L.E. (2018) Developmental temperatures and
- 281 phenotypic plasticity in reptiles: a systematic review and meta-analysis. Biological
- 282 Reviews, **93**, 72–97.
- 283 Parker, T.H., Forstmeier, W., Koricheva, J., Fidler, F., Hadfield, J., En Chee, Y., Kelly,
- 284 C.D., Gurevitch, J. & Nakagawa, S. (2016) Transparency in Ecology and Evolution:
- Real Problems, Real Solutions. Trends in Ecology and Evolution, 31, 711–719.
- Peng, R.D. (2011) Reproducible research in computational science. Science, 334, 1226.
- 287 Peng, R.D., Dominici, F. & Zeger, S.L. (2006) Reproducible epidemiologic research.
- American Journal of Epidemiology, **163**, 783–789.
- 289 Poisot, T. (2011) The digitize package: extracting numerical data from scatterplots.
- 290 The R Journal, **3**, 25–26.
- 291 Rohatgi, A. (2017) WebPlotDigitizer Software, Version 4.0. Austin, Texas, USA.
- 292 Tummers, B. (2006) DataThief Software, Version 3.0.
- 293 Viechtbauer, W. (2010) Conducting Meta-Analyses in R with the metafor Package.
- 294 Journal of Statistical Software, **36**, 1–48.
- 295 Wan, X., Wang, W., Liu, J. & Tong, T. (2014) Estimating the sample mean and
- standard deviation from the sample size, median, range and/or interquartile range.
- 297 BMC Medical Research Methodology, 14, 135.

Figures

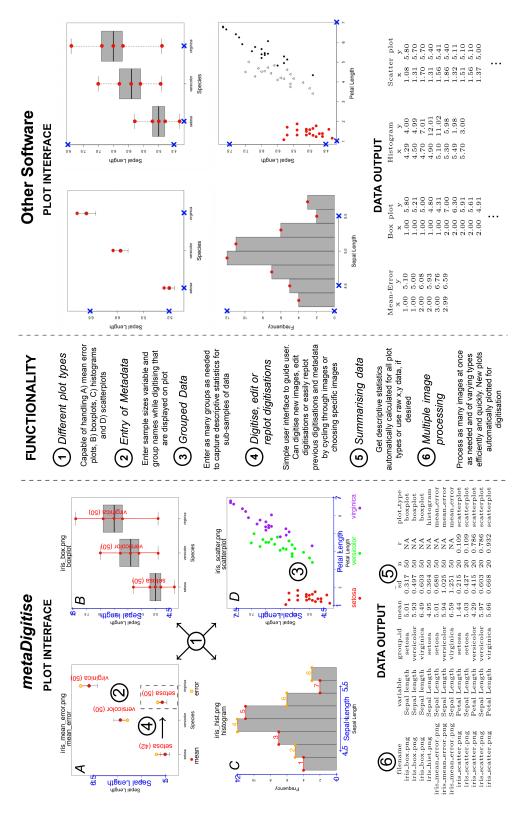


Figure 1: Functionality of **metaDigitise**. Using the iris dataset in R, digitisation of different plot types, A) mean/error plot, B) box plot, C) histogram and D) scatter plot, is shown in **metaDigitise** (left) compared with other common softwares (right). A) and B) are plotted with the whole dataset, C) is just the data for the species *setosa* and D) a subset from all three species. Notable functions of metaDigitise are listed in the center. Other software also perform points 3 and 4 (see Table 1), although these functions are more developed in **metaDigitise**. As shown on the left hand side of the figure, **metaDigitise** clearly displays the stages of the digitisation to aid the transparency of the process, and returns concatenated summary data for all images.

Tables 299

Function	metaDigitise	$GraphClick^1$	$DataThief^2$	$DigitizeIt^3$	$WebPlotDigitizer^4$	$\mathrm{metagear}^5$	digitize ⁶
Scatterplots	>	>	>	>	>	7.	>
Mean/error plots	>	>	>	×	×	~	×
Boxplots	>	×	×	×	×	×	×
Histograms	>	×	×	×	7.7	×	×
Entry of metadata	>	×	×	×	×	×	×
Grouped Data	>	>	×	>	>	×	×
${ m Reproducable}^8$	>	>	>	×	>	>	×
Summarising data	>	×	×	×	×	×	×
Multiple image processing	>	×	×	×	×	×	×
Automated point detection	×	>	×	>	>	>	×
Line extraction	×	>	>	>	>	×	×
Zoom	×	>	>	>	>	×	×
Graph rotation ⁹	>	>	>	>	>	×	×
Log axis	>	>	>	>	>	×	×
Dates	×	×	>	×	>	×	×
Asymmetric error bars	×	×	>	×	×	×	×
Freeware	\checkmark 10	\checkmark^{11}	\checkmark^{11}	\times^{11}	\checkmark^{11}	\checkmark 10	\checkmark 10
1 A wise & Coffee (9000) 2 Thurston	(9006)	3 Demogram (9	(9019) 4 Delegan	. (9017) 5 T	::::::::::::::::::::::::::::::::::::::	:0011)	

 $^{^{1}}$ Arizona-Software (2008) 2 Tummers (2006) 3 Bormann (2012) 4 Rohatgi (2017) 5 Lajeunesse (2016) 6 Poisot (2011)

Table 1: Comparison of functionality between different digitisation softwares.

Only automated, no manual extraction.
 Allows saving, re-plotting and editing of data extraction.
 Or handles rotated graphs.
 R package.
 Standalone software.