

1 **Automated Residual Plot Assessment with the R Package**
2 **autovi and the Shiny App autovi.web**

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Summary

7 Visual assessment of residual plots is a common approach for diagnosing linear
models, but it relies on manual evaluation, which does not scale well and can lead
to inconsistent decisions across analysts. The lineup protocol, which embeds the
observed plot among null plots, can reduce subjectivity but requires even more
human effort. In today's data-driven world, such tasks are well-suited for automation.
We present a new R package that uses a computer vision model to automate the
evaluation of residual plots. An accompanying Shiny app is provided for ease of use.
Given a sample of residuals, the model predicts a visual signal strength (VSS) and
offers supporting information to help analysts assess the adequacy of their model
fit.

8 *Key words:* initial data analysis; statistical graphics; data visualization; visual inference;
computer vision; machine learning; hypothesis testing; regression analysis;
model diagnostics

9 1. Introduction

10 Regression analysis is a widely used statistical modeling technique for data in many
11 fields. There is a vast array of software for conducting regression modeling and
12 generating diagnostics. The package `lmtest` (Zeileis & Hothorn 2002) provides a suite
13 of conventional tests, while the `stats` package (R Core Team 2022) offers standard
14 diagnostic plots such as residuals vs. fitted values, quantile-quantile (Q-Q) plots,
15 and residuals vs. leverage plots. Additional packages like `jtools` (Long 2022), `olsrr`

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16 ([Hebbali 2024](#)), `rockchalk` ([Johnson 2022](#)), and `ggResidpanel` ([Goode & Rey 2019](#))
17 deliver similar graphical diagnostics, often with enhanced aesthetics or interactive
18 features. These tools collectively produce the core diagnostic plots outlined in the
19 classical text by [Cook & Weisberg \(1982\)](#). The `ecostats` package ([Warton 2023](#))
20 extends these diagnostics by incorporating simulation envelopes into residual plots.
21 Meanwhile, `DHARMa` ([Hartig 2022](#)) compares empirical quantiles (0.25, 0.5, and 0.75)
22 of scaled residuals to their theoretical counterparts, with a strong focus on identifying
23 model violations such as heteroscedasticity, misspecified functional forms, and issues
24 specific to generalized linear and mixed-effect models, like over/under-dispersion. It
25 also provides conventional test annotations to reduce the risk of misinterpretation.

26 However, relying solely on subjective assessments of these plots can lead to issues such as
27 over-interpreting random patterns as model violations. [Li et al. \(2024a\)](#) demonstrated
28 that visual methods using the lineup protocol ([Buja et al. 2009](#)) for assessing residuals
29 are more useful, and also perform more practically than conventional tests due to their
30 reduced sensitivity to minor departures. Packages such as `nullabor` ([Wickham et al.
31 2020](#)), `HLMdiag` ([Loy & Hofmann 2014](#)), and `regressinator` ([Reinhart 2024](#)), enable
32 users to compare observed residual plots with plots of samples from null distributions,
33 helping to quantify the significance of any detected patterns.

34 However, as discussed in [Li et al. \(2024b\)](#), the lineup protocol has significant limitations
35 in large-scale applications due to dependence on human labor. Thus, a computer vision
36 model was developed with an associated statistical testing procedure to automate
37 the assessment of residual plots. This model takes a residual plot and a vector of
38 auxiliary variables (such as the number of observations) as inputs and outputs the
39 predicted visual signal strength (VSS). This strength estimates the distance between
40 the residual distribution of the fitted regression model and the reference distribution
41 assumed under correct model specification.

42 To make the statistical testing procedure and trained computer vision model widely
43 accessible, we developed the R package `autovi`, and a web interface, `autovi.web` to
44 make it easy for users to automatically read their residual plots with the trained
45 computer vision model.

46 The remainder of this paper is structured as follows: Section 2 introduces the definition
47 and computation of visual signal strength. Section 3 provides a detailed documentation
48 of the `autovi` package, including its usage and infrastructure. Section 4 focuses on the
49 `autovi.web` interface, describing its design and usage, along with illustrative examples.
50 Finally, Section 5 presents the main conclusions of this work.

51 **2. Definition and computation of visual signal strength**

- 52 To train a computer vision model, a measure of the visible pattern in a plot is needed.
 53 We call this the **visual signal strength** (VSS), which measures how prominently a
 54 specific set of visual patterns appears in an image. This can be computed for a training
 55 set of data, and plots, where the generating distributions are specified.
- 56 In the context of regression model diagnostics, VSS describes the clarity of visual
 57 patterns on a diagnostic plot that may indicate model violations. Violations can be
 58 categorized as weak, moderate, or strong, but here we treat it as a continuous positive
 59 real variable. Importantly, its interpretation depends on how it is linked to a function
 60 of the data or the underlying data generating process. Consequently, the calculation
 61 of VSS can vary across different model classes or within the same model, depending
 62 on the generating function.
- 63 VSS is an estimate of the distance between the residual distribution of a fitted classical
 64 normal linear regression model and a reference distribution; more details can be found
 65 in [Li et al. \(2024b\)](#). The distance measure is based on the Kullback-Leibler (KL)
 66 divergence:

$$D = \log(1 + D_{KL}),$$

67 where D_{KL} is given by:

$$D_{KL} = \int_{\mathbb{R}^n} \log \frac{p(\mathbf{e})}{q(\mathbf{e})} p(\mathbf{e}) d\mathbf{e}, \quad (1)$$

- 68 here, $p(\cdot)$ and $q(\cdot)$ are the probability density functions of the reference residual
 69 distribution P and the true residual distribution Q , respectively.
- 70 This distance measure depends on knowledge of the true residual distribution, which
 71 is unknown in practice. To compute D_{KL} for the training samples, Equation 1 takes
 72 different forms depending on the specific model violations. For instance, where necessary
 73 higher-order predictors, \mathbf{Z} , and their corresponding parameter, $\boldsymbol{\beta}_Z$, are omitted from
 74 the fitted linear model, the distance measure can be expanded as follows:

$$D_{KL} = \frac{1}{2} (\boldsymbol{\mu}_z^\top (\text{diag}(\mathbf{R}\sigma^2))^{-1} \boldsymbol{\mu}_z),$$

75 where $\mu_z = \mathbf{R}\mathbf{Z}\beta_z$, $\mathbf{R} = \mathbf{I}_n - \mathbf{X}(\mathbf{X}^\top \mathbf{X})^{-1}\mathbf{X}^\top$ and \mathbf{X} is the design matrix of the
76 regression model.

77 The computer vision model approximates this mapping from a set of residuals to its
78 corresponding distance measure. It is trained on a large number of synthetic regression
79 models, where the data-generating process is known, allowing the distance measure
80 to be explicitly calculated. The model takes a residual plot as input and outputs the
81 corresponding distance measure. Additional details are provided in [Li et al. \(2024b\)](#).

82 3. R package: autovi

83 The main purpose of **autovi** is to provide rejection decisions and *p*-values for testing
84 the null hypothesis (H_0) that the regression model is correctly specified. The package
85 provides automated interpretation of residual plots using computer vision. The name
86 **autovi** stands for **a**utomated **v**isual **i**nference. This functionality can be accessed
87 through the R package **autovi**, or through a web interface, **autovi.web**, which allows
88 use without the full installation of R, Python, and package dependencies on the user's
89 system. locally.

90 3.1. Motivation

91 Figure 1 shows three sets of plots of residuals against fitted values. The simulated
92 example in (a) might be interpreted as a heteroscedastic pattern, however the
93 automated reading would predict this to have a visual signal strength (VSS) of
94 1.53, with a corresponding *p*-value of 0.25. This means it would be interpreted as
95 a good residual plot, that there is nothing in the data to indicate a violation of
96 model assumptions. Skewness in the predictor variables is generating the apparent
97 heteroscedasticity, where the smaller variance in residuals at larger fitted values is
98 due to smaller sample size only. The Breusch-Pagan test ([Breusch & Pagan 1979](#)) for
99 heteroscedasticity would also not reject this as good residual plot.

100 The data in (b) is generated by fitting a linear model predicting `mpg` based on `hp`
101 using the datasets::`mtcars`. It is a small data set, and there is a hint of nonlinear
102 structure not captured by the model. The automated plot reading would predict a
103 VSS of 3.57, which has a *p*-value less than 0.05. That is, the nonlinear structure is
104 most likely real, and indicates a problem with the model. The conventional test, a
105 Ramsey Regression Equation Specification Error Test (RESET) ([Ramsey 1969](#)) would
106 also strongly detect the nonlinearity.

107 The third example is generated using the `surreal` package (Balamuta 2024) where
 108 structured residuals are hidden in data, to be revealed if the correct model is specified.
 109 Here a quote based on Tukey is used as the residual structure “visual summaries focus
 110 on unexpected values”. The automated plot reading predicts the VSS to be 5.87, with
 111 a p -value less than 0.05. This structure is blindingly obvious visually, but a RESET
 112 test for nonlinear structure would not report a problem. (It would be detected by
 113 a Breusch-Pagan for heteroscedasticity and also Shapiro-Wilk test (Shapiro & Wilk
 114 1965) for non-normality.)

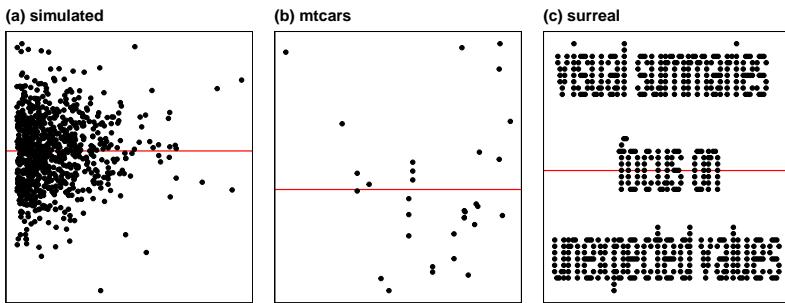


Figure 1. Reading residual plots can be a difficult task, particularly for students new to statistical modeling. The `autovi` package makes it easier. Here are three examples of residual plots, which may appear to have structure. According to `autovi`, the visual signal strengths (VSS) of these three examples are approximately (a) 1.53, (b) 3.57, (c) 5.87, resulting in (b), (c) being significant violations of good residuals, but (a) is consistent with a good residual plot.

115 3.2. Implementation

116 The `autovi` package is built on the `bandicoot` object-oriented programming (OOP)
 117 system (Li 2024), marking a departure from R’s traditional S3 generic system. This
 118 OOP architecture enhances flexibility and modularity, allowing users to redefine key
 119 functions through method overriding.

120 The `autovi` infrastructure effectively integrates multiple programming languages and
 121 libraries into a comprehensive analytical tool. It relies on five core libraries from
 122 Python and R, each playing a critical role in the analysis pipeline. In Python, `pillow`
 123 (Clark et al. 2015) handles image processing tasks such as reading and resizing PNG
 124 files of residual plots, then converting them into input tensors for further analysis.
 125 `TensorFlow` (Abadi et al. 2016), a key component of modern machine learning, is used
 126 to predict the VSS of these plots using a pre-trained convolutional neural network.

127 In the R environment, `autovi` utilizes several libraries. `ggplot2` (Wickham 2016)
 128 generates the initial residual plots, saved as PNG files for visual input. `cassowaryr`
 129 (Mason et al. 2022) computes scagnostics (scatter plot diagnostics), providing numerical
 130 features that capture statistical properties of the plots. These scagnostics complement
 131 the visual analysis by offering quantitative metrics as secondary input to the
 132 computer vision model. `reticulate` (Ushey, Allaire & Tang 2024) enables seamless
 133 communication between R and Python.

134 **3.3. Installation**

135 The `autovi` package is available on CRAN. It is actively developed and maintained,
 136 with the latest updates accessible on GitHub. This paper uses `autovi` version 0.4.1.
 137 The package includes internal functions to check the current Python environment used
 138 by the `reticulate` package. If the necessary Python packages are not installed in the
 139 Python interpreter, an error will be raised. If you want to select a specific Python
 140 environment, you can do so by calling the `reticulate::use_python()` function before
 141 using the `autovi` package.

142 We recommend using the Shiny app `autovi.web` if users encounter installation
 143 problems.

144 **3.4. Usage**

145 **3.4.1. Numerical summary**

146 Three steps are needed to get an automated assessment of a set of residuals and fitted
 147 values:

- 148 1. Load the `autovi` package using the `library()` function.
- 149 2. Create a checker object with a linear regression model.
- 150 3. Call the `check()` method of the checker, which, by default, predicts the VSS for
 the true residual plot, 100 null plots, and 100 bootstrapped plots. The method
 stores the predictions internally and prints a concise results report.

153 The code to do this is:

```
library(autovi)
checker <- residual_checker(lm(dist ~ speed, data = cars))
checker$check()
```

154 It produces the following summary:

155

```

156 -- <AUTO_VI object>
157 Status:
158 - Fitted model: lm
159 - Keras model: UNKNOWN
160 - Output node index: 1
161 - Result:
162 - Observed visual signal strength: 3.162 (p-value = 0.0396)
163 - Null visual signal strength: [100 draws]
164 - Mean: 1.274
165 - Quantiles:

166
167      25%    50%    75%    80%    90%    95%    99%
168      0.8021  1.1109  1.5751  1.6656  1.9199  2.6564  3.3491
169

170 - Bootstrapped visual signal strength: [100 draws]
171 - Mean: 2.786 (p-value = 0.05941)
172 - Quantiles:

173
174      25%    50%    75%    80%    90%    95%    99%
175      2.452   2.925   3.173   3.285   3.463   3.505   3.652
176

177 - Likelihood ratio: 0.7275 (boot) / 0.06298 (null) = 11.55

```

178 The summary includes observed VSS of the true residual plot and associated *p*-value
 179 of the automated visual test. The *p*-value is the proportion of null plots (out of the
 180 total 100) that have VSS greater than or equal to that of the true residual plot. The
 181 report also provides sample quantiles of VSS for null samples and bootstrapped data
 182 plots, providing more information about the sampling variability and a likelihood of
 183 model violations. The likelihood is computed from the proportion of values greater
 184 than the observed VSS in both the bootstrapped data values and the simulated null
 185 values.

186 **3.4.2. Visual summary**

187 Users can visually inspect the original residual plot alongside a sample null plot using
 188 `plot_pair()` or a lineup of null plot `plot_lineup()`. This visual comparison can
 189 clarify why H_0 is either rejected or not, and help identify potential remedies.

```
checker$plot_pair()
```

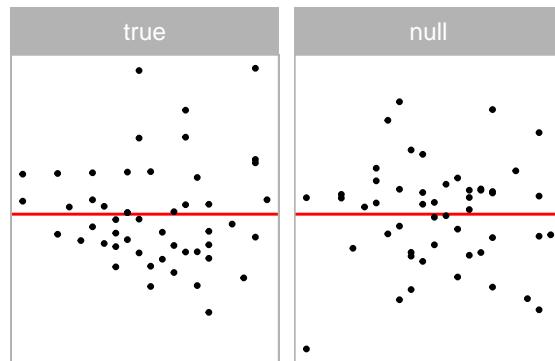


Figure 2. True plot alongside one null plot, for quick comparison.

190 The `plot_pair()` method (Figure 2) displays the true residual plot on the left and a
 191 single null plot on the right. If a full lineup was shown, the true residual plot would
 192 be embedded in a page of null plots. Users should look for any distinct visual patterns
 193 in the true residual plot that are absent in the null plot. Running these functions
 194 multiple times can help any visual suspicions, as each execution generates new random
 195 null plots for comparison.

196 The package offers a straightforward visualization of the assessment result through
 197 the `summary_plot()` function.

```
checker$summary_plot()
```

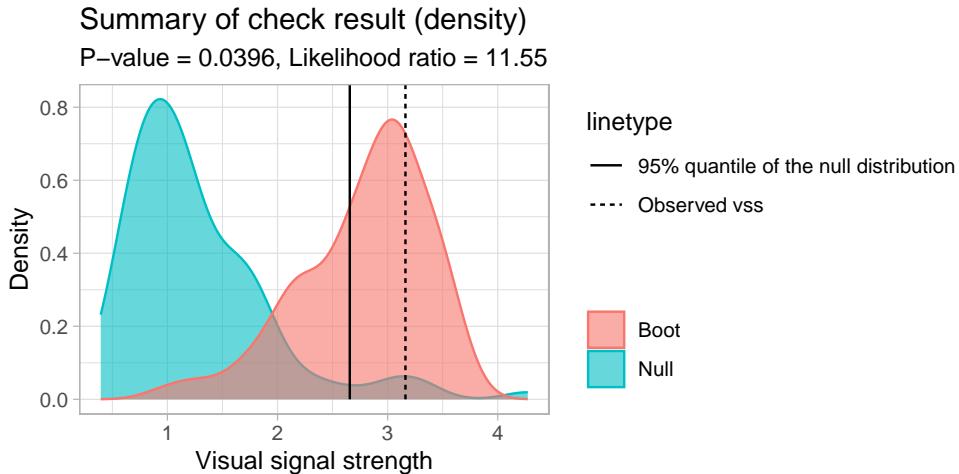


Figure 3. Summary plot comparing the densities of VSS for bootstrapped residual samples (red) relative to VSS for null plots (blue).

- 198 In the result, shown in Figure 3, the blue area represents the density of VSS for null
 199 residual plots, while the red area shows the density for bootstrapped residual plots.
 200 The dashed line indicates the VSS of the true residual plot, and the solid line marks
 201 the critical value at a 95% significance level. The p -value and the likelihood ratio are
 202 displayed in the subtitle. The likelihood ratio represents the ratio of the likelihood
 203 of observing the VSS of the true residual plot from the bootstrapped distribution
 204 compared to the null distribution.
- 205 Interpreting the plot involves several key aspects. If the dashed line falls to the right of
 206 the solid line, it suggests rejecting the null hypothesis. The degree of overlap between
 207 the red and blue areas indicates similarity between the true residual plot and null
 208 plots; greater overlap suggests more similarity. Lastly, the portion of the red area to
 209 the right of the solid line represents the percentage of bootstrapped models considered
 210 to have model violations.
- 211 This visual summary provides an intuitive way to assess the model's fit and potential
 212 violations, allowing users to quickly grasp the results of the automated analysis.

213 3.5. Modularized infrastructure

- 214 The initial motivation for developing `autovi` was to create a convenient interface for
 215 sharing the models described and trained in Li et al. (2024b). However, recognizing
 216 that the classical normal linear regression model represents a restricted class of

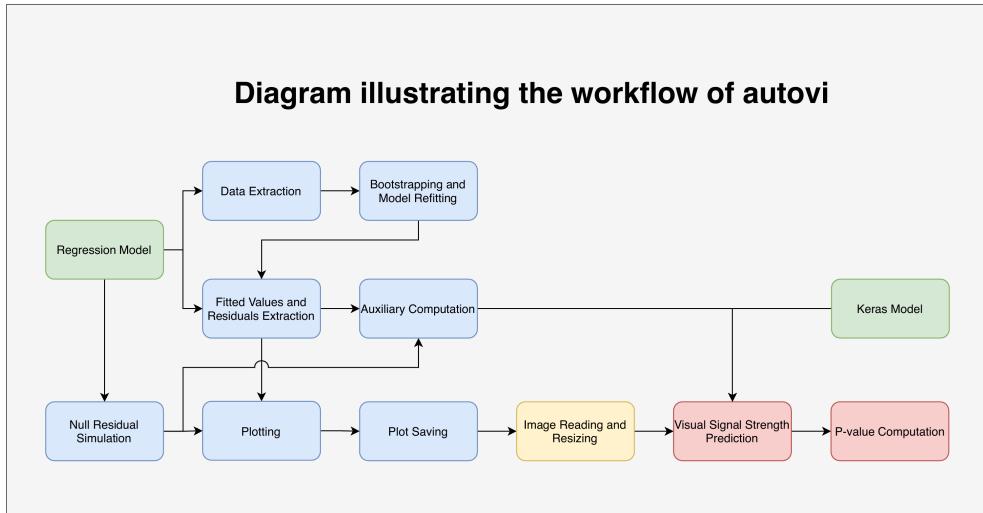


Figure 4. Diagram illustrating the infrastructure of the R package `autovi`. The modules in green are primary inputs provided by users. Modules in blue are overridable methods that can be modified to accommodate users' specific needs. The module in yellow is a pre-defined non-overridable method. The modules in red are primary outputs of the package.

models, we sought to avoid limiting the potential for future extensions, whether by the original developers or other developers. As a result, the package was designed to function seamlessly with linear regression models with minimal modification and few required arguments, while also accommodating other classes of models through partial infrastructure substitution. This modular and customizable design allows `autovi` to handle a wide range of residual diagnostics tasks.

The infrastructure of `autovi` consists of ten core modules: data extraction, bootstrapping and model refitting, fitted values and residuals extraction, auxiliary computation, null residual simulation, plotting, plot saving, image reading and resizing, VSS prediction, and *p*-value computation. Each module is designed with minimal dependency on the preceding modules, allowing users to customize parts of the infrastructure without affecting its overall integrity. An overview of this infrastructure is illustrated in Figure 4.

The modules for VSS prediction and *p*-value computation are predefined and cannot be overridden, although users can interact with them directly through function arguments. Similarly, the image reading and resizing module is fixed but will adapt to different Keras models by checking their input shapes. The remaining seven modules are designed to be overridable, enabling users to tailor the infrastructure to their specific needs. These modules are discussed in detail in the package documentation.

236

4. Web interface: `autovi.web`

237 The `autovi.web` shiny application extends the functionality of `autovi` by offering a
238 user-friendly web interface for automated residual plot assessment. This eliminates the
239 common challenges associated with software installation, so users can avoid managing
240 Python environments or handling version requirements for R libraries. The platform
241 is cross-platform and accessible on various devices and operating systems, making it
242 suitable even for users without R programming experience. Additionally, updates are
243 managed centrally, ensuring that users always have access to the latest features. This
244 section discusses the implementation based on `autovi.web` version 0.1.0.

245

4.1. Implementation

246 The interface `autovi.web` is built using the `shiny` (Chang et al. 2022) and
247 `shinydashboard` (Chang & Borges Ribeiro 2021) R packages. Hosted on the
248 `shinyapps.io` domain, the application is accessible through any modern web browser.
249 The R packages `htmltools` (Cheng et al. 2024) and `shinycssloaders` (Sali & Attali
250 2020) are used to render markdown documentation in shiny application, and for loading
251 animations for shiny widgets, respectively.

252 Determining the best way to implement the backend was difficult. In our initial
253 planning for `autovi.web`, we considered implementing the entire web application using
254 the `webr` framework (Moon 2020), which would have allowed the entire application to
255 run directly in the user’s browser. However, `webr` does not support packages which use
256 compiled fortran code, which is required by `splancs` (Rowlingson & Diggle 2023), a
257 dependency of `autovi`. In the future, it is possible that a working Emscripten (Zakai
258 2011) version of this package may allow full `webr` support.

259 We also explored the possibility of implementing the web interface using frameworks
260 built on other languages, such as Python. However, server hosting domains that
261 natively support Python servers typically do not have the latest version of R installed.
262 Additionally, calling R from Python is typically done using the `rpy2` Python library
263 (Gautier 2024), but this approach can be awkward when dealing with language syntax
264 related to non-standard evaluation. Another option we considered was renting a server
265 where we could have full control, such as those provided by cloud platforms like
266 Google Cloud Platform (GCP) or Amazon Web Services (AWS). However, deploying
267 and maintaining the server securely requires some expertise. Ultimately, the most
268 practical solution was to use the `shiny` and `shinydashboard` frameworks, which are

269 well-established in the R community and offer a solid foundation for web application
270 development.

271 The server-side configuration of `autovi.web` is carefully designed to support its
272 functionality. Most required Python libraries, including `pillow` and `numpy`, are pre-
273 installed on the server. These libraries are integrated into the Shiny application using
274 the `reticulate` package, which provides an interface between R and Python.

275 Due to the resource allocation policy of shinyapps.io, the server enters a sleep mode
276 during periods of inactivity, resulting in the clearing of the local Python virtual
277 environment. Consequently, when the application “wakes up” for a new user session,
278 these libraries need to be reinstalled. While this ensures a clean environment for each
279 session, it may lead to slightly longer loading times for the first user after a period of
280 inactivity.

281 In contrast to `autovi`, `autovi.web` leverages `TensorFlow.js`, a JavaScript library
282 that allows the execution of machine learning models directly in the browser. This
283 choice enables native browser execution, enhancing compatibility across different user
284 environments, and shifts the computational load from the server to the client-side.
285 `TensorFlow.js` also offers better scalability and performance, especially when dealing
286 with resource-intensive computer vision models on the web.

287 While `autovi` requires downloading the pre-trained computer vision models from
288 GitHub, these models in “.keras” file format are incompatible with `TensorFlow.js`.
289 Therefore, we extract and store the model weights in JSON files and include
290 them as extra resources in the Shiny application. When the application initializes,
291 `TensorFlow.js` rebuilds the computer vision model using these pre-stored weights.

292 To allow communication between `TensorFlow.js` and other components of the Shiny
293 application, the `shinyjs` R package ([Attali 2021](#)) is used. This package allows calling
294 custom JavaScript code within the Shiny framework. The specialized JavaScript
295 code for initializing `TensorFlow.js` and calling `TensorFlow.js` for VSS prediction is
296 deployed alongside the Shiny application as additional resources.

297 4.2. Usage

298 The workflow of `autovi.web` is designed to be straightforward, with numbered
299 steps displayed in each panel. There are two example datasets provided by the
300 web application. The single residual plot example uses the `dino` dataset from the
301 R package `datasauRus` ([Davies, Locke & D'Agostino McGowan 2022](#)). The lineup

example uses residuals from a simulated regression model that has a non-linearity issue. We walk through the lineup example to further demonstrate the workflow of the web application.

4.2.1. Reading data and setting parameters

The user can select to upload data as either a single set of residuals and fitted values in a two (or more) column CSV file or a pre-computed lineup of residuals and null datasets in a three (or more) column CSV file (i.e. multiple sets of residuals and fitted values with a column indicating the set label). Here we illustrate use with lineup example data sets (Figure 5). To use the lineup example data, click the “Use Lineup Example” button. The data status will then update to show the number of rows and columns in the dataset, and the CSV type will automatically be selected to the correct option. Since the example dataset follows the variable naming conventions assumed by the web application, the columns for fitted values, residuals, and labels of residual plots are automatically mapped such that the column named as `.fitted` is mapped to fitted values, `.resid` is mapped to residuals and if applicable, `.sample` to labels of the residual set (middle image). If the user is working with a custom dataset, these options must be set accordingly. Whenever a data containing a lineup, the user must manually select the label for the true residual plot, otherwise the web application does not provide all the results. The last step is to click the play button (right image) to start the assessment.

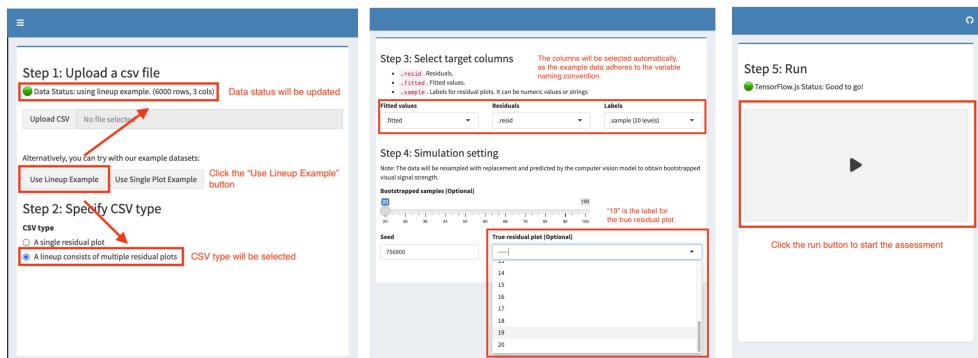


Figure 5. To begin the workflow for `autovi` using the lineup example dataset, the user clicks the “Use Lineup Example” button (left) to load the example dataset, during which the data status and CSV type will be automatically updated. The user must manually select the label for the true residual plot (middle) to compute further results. The user initiates the assessment of the lineup example data by clicking the run button (right).

322 **4.2.2. Results provided**

323 Results are provided in multiple panels. The first row of the table Figure 6 is the most
 324 crucial to check, as it provides the VSS and the rank of the true residual plot among
 325 the other plots. The summary text beneath the table provides the *p*-value, which can
 326 be used for quick decision-making. The lineup is for manual inspection, and the user
 327 should see if the true residual plot is visually distinguishable from the other plots, to
 328 confirm if the model violation is serious.

329 The density plot in Figure 7 offers a more robust result, allowing the user to compare
 330 the distribution of bootstrapped VSS with the distribution of null VSS. Finally, the
 331 grayscale attention map (right image) can be used to check if the target visual features,
 332 like the non-linearity present in the lineup example, are captured by the computer
 333 vision model, ensuring the quality of the assessment.

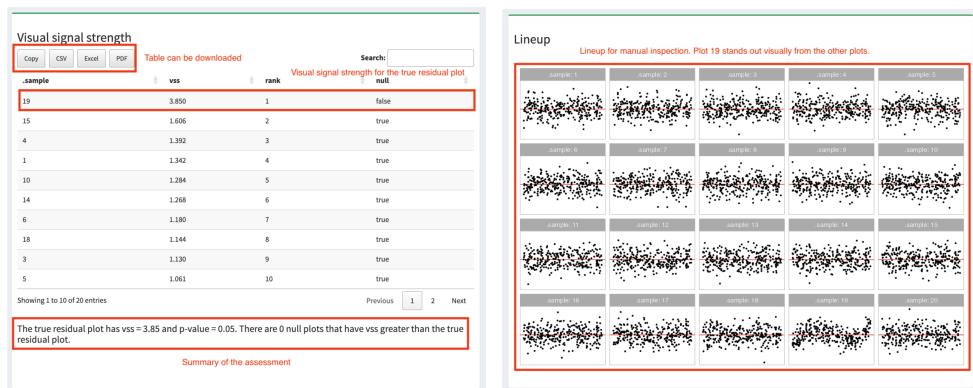


Figure 6. Results for the lineup. The VSS of the true residual plot is displayed in the first row of the table of VSS values for all the null plots (left image), with a summary text beneath the table providing the *p*-value to aid in decision-making. A lineup of residual plots allows for manual inspection (right image).

334

5. Conclusions

335 This paper presents new regression diagnostics software, the R package **autovi** and
 336 its accompanying web interface, **autovi.web**. It addresses a critical gap in the current
 337 landscape of statistical software. While regression tools are widely available, effective
 338 and efficient diagnostic methods have lagged behind, particularly in the field of residual
 339 plot interpretation.

340 The **autovi** R package, introduced in this paper, automates the assessment of residual
 341 plots by incorporating a computer vision model, reducing reliance on time-consuming

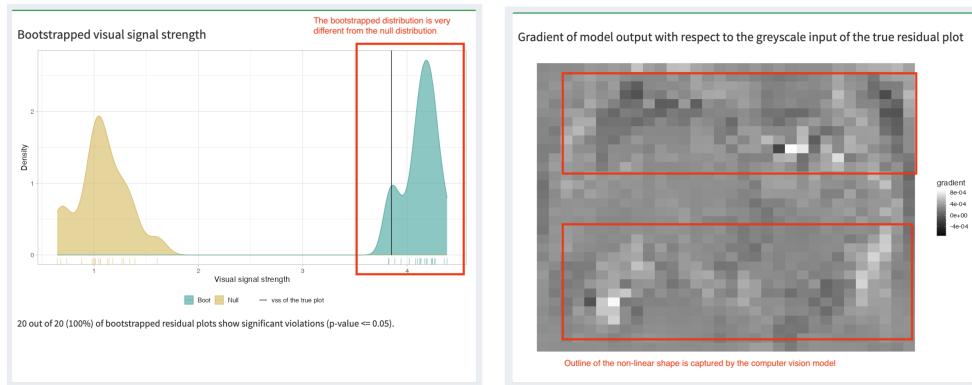


Figure 7. Summaries assessing the strength of the pattern and which elements of the plot contribute. The density plot helps verify if the bootstrapped distribution differs from the null distribution (left image). The attention map (right image) offers insights into whether the computer vision model has captured the intended visual features of the true residual plot.

342 and potentially inconsistent human interpretation. This automation improves the
 343 efficiency of the diagnostic process and promotes consistency in model evaluation
 344 across different users and studies.

345 The development of the accompanying Shiny app, **autovi.web**, expands access to these
 346 advanced diagnostic tools, by providing a user-friendly interface. It makes automated
 347 residual plot assessment accessible to a broader audience, including those who may not
 348 have extensive programming experience. This web-based solution effectively addresses
 349 the potential barriers to adoption, such as complex dependencies and installation
 350 requirements, that are often associated with advanced statistical software.

351 The combination of **autovi** and **autovi.web** offers a comprehensive solution to the
 352 challenges of residual plot interpretation in regression analysis. These tools have the
 353 potential to significantly improve the quality and consistency of model diagnostics
 354 across various fields, from academic research to industry applications. By automating
 355 a critical aspect of model evaluation, they allow researchers and analysts to focus more
 356 on interpreting results and refining models, rather than grappling with the intricacies
 357 of plot assessment.

358 The framework established by **autovi** and **autovi.web** opens up exciting possibilities
 359 for further research and development. Future work could explore the extension of these
 360 automated assessment techniques to other types of diagnostic plots and statistical
 361 models, potentially revolutionizing how we approach statistical inference using visual
 362 displays more broadly.

363 **6. Resources and supplementary material**

364 The current version of `autovi` can be installed from CRAN, and source
 365 code for both packages are available at github.com/TengMCing/autovi and
 366 github.com/TengMCing/autovi_web respectively. The web interface is available from
 367 autoviweb.netlify.app.

368 This paper is reproducibly written using Quarto ([Allaire et al. 2024](#)) powered by
 369 Pandoc ([MacFarlane, Krewinkel & Rosenthal 2024](#)) and pdfTeX. The full source code
 370 to reproduce this paper is available at github.com/TengMCing/autovi_paper.

371 These R packages were used for the work: `tidyverse` ([Wickham et al. 2019](#)), `lmtest`
 372 ([Zeileis & Hothorn 2002](#)), `kableExtra` ([Zhu 2021](#)), `patchwork` ([Pedersen 2022](#)),
 373 `rcartocolor` ([Nowosad 2018](#)), `glue` ([Hester & Bryan 2022](#)), `here` ([Müller 2020](#)),
 374 `magick` ([Ooms 2023](#)), `yardstick` ([Kuhn, Vaughan & Hvitfeldt 2024](#)) and `reticulate`
 375 ([Ushey, Allaire & Tang 2024](#)).

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