

DSAF Shiny Prototype Application Manual

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Summary

The attached DSAF Shiny Application allows users to view the status of a hypothetical fish population – separated between multiple jurisdictional zones in Canada – assessed by six indicators of species' change. To balance computation and interactivity, the app relies on precomputed visualizations, which are uploaded upon loading and displayed according to user inputs.

Here, I outline the contents of the application folder, including scripts to preprocess data, scripts to create interactive visualizations, intermediate data objects, and application scripts.

Directory overview

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└── Data/	
└── Data_SHinyApp_Proof_of_Concept/	Various input data from DSAF team
└── eez_can/	Shapefile of Canadian EEZ
└── shiny/	
└── preprocessing_scripts/	Scripts for making interactive plots
└── unscripted/	Non-R files to generate app objects (logos & pdf)
└── shiny_processed_data/	Intermediate data objects for plots
└── www/	Final objects to load into App
└── ui.R	User interface - overall
└── ui_Tab1.R	Module for Page 1 (About)
└── ui_Tab2.R	Module for Page 2 (Data Page)
└── ui_Tab3.R	Module for Page 3 (Science)
└── ui_Tab4.R	Module for Page 4 (Download)
└── app_text.R	All application text, saved as list
└── server.R	Server logic
└── global.R	Global object processing when application loads

Making / Editing Plots

All plots in the application are pre-rendered from scripts in the “preprocessing_scripts/” folder, and saved into the “www/” folder for use in the Shiny app. In “preprocessing_scripts”, the file “00_plot_opts.R” controls global options (colors to represent each region) that apply to all following plots. The script, “01_preprocess_all.R” sources all other scripts in this folder to create interactive plots that are used in the application. For example, if the variables in “00_plot_opts.R” are changed, running “01_preprocess_all.R” will override the existing plots with the new settings.

Editing Application Text

For organization, all text from the shiny app is stored in the script, “app_text.R”. In this script, all text objects are organized into a hierarchical list object, with sublists for text in each individual application section: header, tab 1, tab 2, and so on. This script is sourced by the application upon loading, and the list is queried within each UI script. Changes here will be reflected when the application is run. Note that some of the elements are character strings, but some are wrapped in html functions (e.g., h2(“text”)) or written with html formatting (e.g., HTML(“some words are bolded”)). New text can either follow this same structure, or be entered directly into the UI pages of the app.

Application Logic

The application logic is simple. The “global.R” script uploads all pre-made plots for a given species (here, “sppx”) into a list, and the server pulls elements from this list when rendering plots according to user inputs. When the user selects “Species X”, the server pulls from the “sppx_plots” list, and when the user queries the abundance plot (“p_abundance”), the server renders an output from sppx_plots\$p_abundance. This structure results in fast rendering and stability, since the computation for plots is done in advance, but it is likely to face limits when scaling up to many species since the application would need to store every plot in advance.

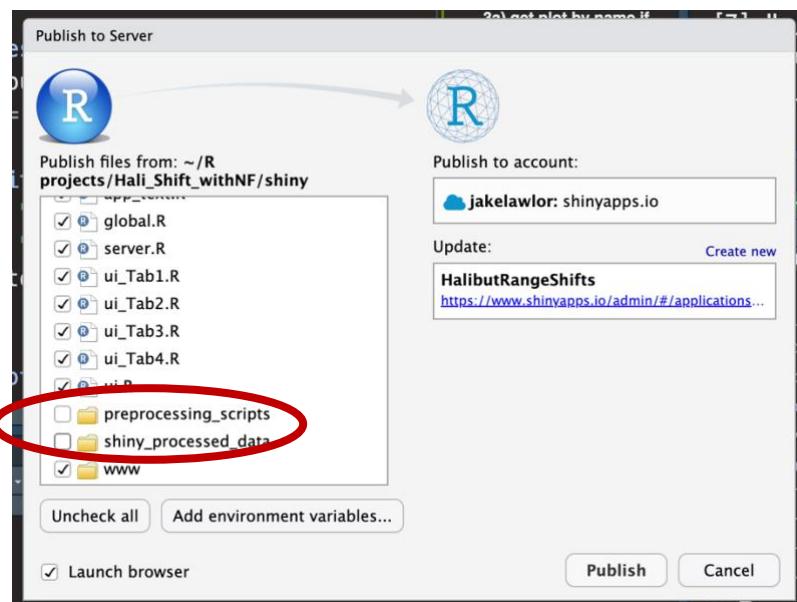
Placeholder Objects

This application also relies on a few pre-rendered objects that are placeholders, not created directly from the provided data. These objects are meant for demonstration purposes only, but could be further developed in the future. For example, the “download” button on UI Tab 4 is meant to show how custom summary outputs could be

programmatically written from application data: when a user selects a focal species and study area, a summary of the species' status in that region could be generated. Here, we use a stored static pdf ("shiny/www/example_summary.pdf") to demonstrate what this might look like. Replace this file to change the downloaded pdf output.

Hosting the Application

Not all files in the "shiny/" folder are necessary for hosting the application on a Shiny server, and excluding unnecessary files will save space and improve efficiency. Inside the "shiny/" folder, "preprocessing_scripts/" and "shiny_preprocessed_data/" do not need to be included in order to publish the application. When publishing from RStudio, these folders can be un-selected in the publish window (see below). For publishing, all app scripts (UI, server, global, and text) are necessary, and the "www/" folder.



Unscripted Files

Some development for this application required files that I created manually outside of R, including custom logos and the downloadable pdf species' summary example. I included these files in the repository in "shiny/preprocessing_scripts/unscripted/". Their outputs were manually saved to the "www/" folder.

Contact

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