How to create a Shiny web app in R

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Outline

- What is Shiny and how to get started?
- ► Simple example
- ► My experience with Shiny
- ▶ Using Shiny as an user interface for an R- package.
- ► How to share your Shiny

What is Shiny?

"A Shiny app is a web page connected to a computer running a live R session" - Shiny cheat sheet

And all you need is:

- ▶ R-studio
- install.packages("shiny") and library(shiny)
- Basic R knowledge (for using templates) and good R knowledge for doing more tailored apps

Templates

https://shiny.rstudio.com/gallery/

Shiny file structure

- ▶ ui.R define the user interface
- ► server.R compute the function, graph etc. depending on the input from the user
- global.R contain all additional code for the app
- ▶ All files are saved in a folder with the name of the app



How to learn Shiny (except for this tutorial)

Basic tutorials

https://shiny.rstudio.com/tutorial/

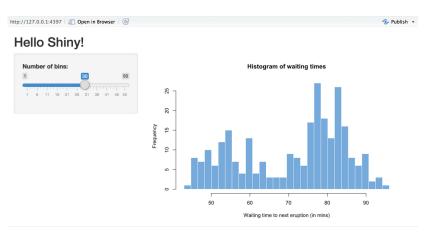
https://www.youtube.com/watch?v=sJl0EE_RE4o&list= PLH6mU1kedUy-aGYi-w1XqSiGtViFK9NpI

https://github.com/aagarw30/R-Shinyapp-Tutorial

Shiny-cheatsheet

http://shiny.rstudio.com/images/shiny-cheatsheet.pdf

Simple example



https://shiny.rstudio.com/articles/basics.html

ui.R

```
# UI for app that draws a histogram
ui <- fluidPage(
# App title ----
                               → Hello Shiny!
  titlePanel("Hello Shiny!"),
# Sidebar layout with input
# and output definitions
                                    Number of bins:
  sidebarLayout(
# Sidebar panel for inputs
   sidebarPanel(
# Input: Number of bins
  sliderInput(inputId = "bins",
      label = "Number of bins:".
      min = 1.
      max = 50,
      value = 30
# Main panel for displaying outputs
   mainPanel(
# Output: Histogram ----
      plotOutput(outputId = "distPlot")
```

ui.R and server.R

```
@ ui.R ×
# UI for app that draws a histogram
ui <- fluidPage(
# App title ----
  titlePanel("Hello Shiny!"),
# Sidebar layout with input
# and output definitions
  sidebarLayout(
# Sidebar panel for inputs
    sidebarPanel(
# Input: Number of bins
  sliderInput(inputId = "bins"
      label = "Number of bins:",
      min = 1.
      max = 50.
      value = 30)
# Main panel for displaying outputs
    mainPanel(
# Output: Histogram ----
      plotOutput(outputId = "distPlot")
```

```
    server.R 

    ≈

# Server logic to draw a histogram
server <- function(input, output){</pre>
output$distPlot <- renderPlot({
     <- faithful$waiting
bins \leftarrow seq(min(x), max(x),
           length.out = input$bins
hist(x, breaks = bins, col = "#75AA
border = "white".
xlab = "Waiting time to next erupti
main = "Histogram of waiting times"
 })
```

My Shiny application

Imagine you want to visualize this:

$$AF(p, b, k, h) = \frac{\Phi\{\Phi^{-1}(p), -b; h\} - \Phi\{\Phi^{-1}(p) - kh, -b; h\}}{p}$$

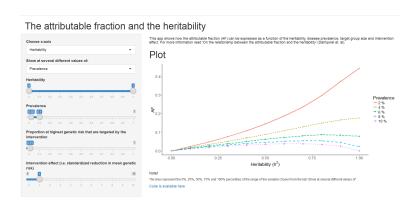
where

 $ightharpoonup \Phi(\cdot)$ is the standard (i.e. mean 0, variance 1) normal distribution function

Problem: A function that depends on 4 parameter is difficult to visualize.

Solution: Interactive graph

My Shiny app



https://afheritability.shinyapps.io/afheritability/

Creating interactive panels

conditionalPanel() looks intuitive since the value choosen in the first condition give you different conditions depending on the value.

My problem: A unique id is needed within every level of the conditionalPanel() to make it interactive \rightarrow was not possible for me

My solution: Create the a function that create sliders depending on the first input arguments.

My solution - ui.R

library(shiny)

```
Only the first
shinvUI(fluidPage(
                                                                   sidebarPanel is made in
  # Application title
 headerPanel("The attributable fraction and the heritability"),
                                                                   ui.R
  sidehar avout (
   SidebarPanel(
  selectInput("xaxis", "Choose x-axis", choices = unique(alternatives$xaxis)).
  selectInput("comparé", "Show at several different values of:", choices= "", selected=""
  uiOutput("Heritability_slider"),
  uiOutput("Prevalence_slider").
                                         The sliders which depend on the value in
  uiOutput("Target_slider"),
                                         the first sidebar will be output in ui.R but
  uiOutput("Intervention slider"
                                         made in server.R
   # Show a plot of the AF and heritability
   mainPanel(h5("This app shows how the attributable fraction (AF) can be expressed as a fun-
             h1("Plot").
     tags style (type="text/css".
                 ".shiny-output-error { visibility: hidden: }".
                 ".shinv-output-error:before { visibility: hidden: }"
     plotOutput("AFfunction").
     h5("Note!").
     h6("The lines represent the 0%, 25%, 50%, 75% and 100% percentiles of the range of the
     helpText( a("Code is available here".
                                              href="https://github.com/ElisabethDahlgwist
```

My solution - server.R

```
shinvServer(function(input, output, session) {
 observeEvent(
   input$xaxis.
 updateSelectInput(session. "compare". "Show at several different values of:".
                  choices = alternatives$compare[alternatives$xaxis == input$xaxis]))
 output in ui.R
   values <- value_maker(input$xaxis, input$compare)</pre>
                  Determine which variable is fixed depending on first input
   if(length(values$H_value)==1) sliderInput("Heritability", "Heritability", min=0, max = 1
   else sliderInput("Heritability", "Heritability", min=0, max = 1, value = c(values$H valu
 output$Prevalence slider <- renderUI({
   values <- value maker(input$xaxis. input$compare)</pre>
   if(length(values$P value)==1) sliderInput("Prevalence", "Prevalence", min=0, max = 1, va
   else sliderInput("Prevalence", "Prevalence", min=0, max = 1, value = c(values$P_value[1]
 output$Target slider <- renderUI({
   values <- value maker(input$xaxis, input$compare)</pre>
```

My solution - global.R

```
A A .
######### Functions for shiny app
alternatives <- data.frame(xaxis = c("Heritability", "Heritability", "Heritability", "Preva
                  compare = c("Prevalence", "Target", "Intervention", "Heritability", "Target"
                  row.names = NULL, stringsAsFactors = FALSE) All combinations of
value_maker <- function(xaxis, compare){</pre>
  if(xaxis == "Heritability" && compare == "Prevalence") {
                                                             "compare".
    H value <- c(0, 1)
                              Heritability and prevalence are allowed to vary but
    P_{value} <- c(0.02.0.1)
    T value <- 0.01
                              not target or intervention.
    T_value <- 1
  if(xaxis == "Heritability" && compare == "Target") {
   H value \leftarrow c(0. 1)
    P value <- 0.5
    T_value <- c(0.01, 0.3)
    T value <- 1
  if(xaxis == "Heritability" && compare == "Intervention") {
    H value <- c(0.1)
    P value <- 0.3
    T_value <- 0.05
    I value <-c(1.5)
  if(xaxis == "Prevalence" && compare == "Heritability") {
    H_value <- c(0.2, 0.6)
    P_value <- c(0, 1)
    T_value <- 0.05
    I value <- 1
```

The render function

Outputs - render*() and *Output() functions work together to add R output to the UI



DT::renderDataTable(expr. options, callback, escape, env, quoted)



dataTableOutput(outputId, icon, ...)



renderImage(expr, env, quoted, deleteFile)



renderPlot(expr. width, height, res, ..., env. auoted, func)



renderPrint(expr. env. auoted, func. width)



renderTable(expr,..., env, quoted, func)



renderText(expr, env, quoted, func)



renderUI(expr, env, quoted, func)

imageOutput(outputId, width, height, click, dblclick, hover, hoverDelay, hoverDelayType, brush, clickld, hoverld, inline)

plotOutput(outputId, width, height, click, dblclick, hover, hoverDelay, hoverDelayType. brush, clickId, hoverId, inline)

verbatimTextOutput(outputId)

tableOutput(outputId)

textOutput(outputId, container, inline)

uiOutput(outputId, inline, container, ...) & htmlOutput(outputId, inline, container, ...)



Shiny as an interface for an R-package

EpiEstim: a package to estimate disease transmissibility during an infectious disease outbreak

Info: https://github.com/jstockwin/EpiEstimApp/wiki

```
install.packages("devtools")
library(devtools)
devtools::install_github("jstockwin/EpiEstimApp",
  ref = "recon-update", force = TRUE)
EpiEstimApp::runEpiEstimApp()
```

Share your Shiny!

There are several alternatives in how to share your Shiny app: https:

//shiny.rstudio.com/articles/deployment-web.html

- Code available on GitHub require that the user have R installed
- Shinyapps.io you share your code with Shinyapps.io. Different alternatives for scaling.
- Shiny Server open-source software requires a Linux server that you will need to set up and maintain
- ► Shiny Server Pro extension of Shiny Server with higher security, control, and support that workgroups and enterprises need

Publish the Shiny app at Shinyapps.io

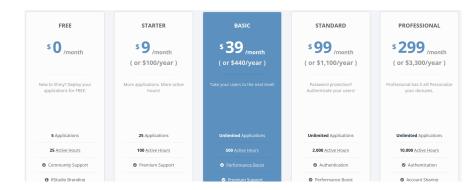
Where to create an account: http://www.shinyapps.io/ How to do it:

https://shiny.rstudio.com/articles/shinyapps.html

Steps to make it happen!!

- You need: install.packages('rsconnect') and library(rsconnect)
- ► Create a Shinyapps.io account
- Configure rsconnect:
 rsconnect::setAccountInfo(name="<ACCOUNT>",
 token="<TOKEN>", secret="<SECRET>")
- ► PUBLISH: rsconnect::deployApp('Directory of your Shiny app folder')

Prices at shinyapps.io



My experience of Shiny in summary

- ► Easy to get started
- With more advanced operations it can be a bit tricky
- BUT, information, tutorials etc. are really really good!
- Surprizingly easy to make the Shiny into a web app but probably more complicated for some more advanced Shiny

My grade to Shiny as a very beginner:



Thank you!

