



Interactive Plotting Libraries - 1

One should look for what is and not what he thinks should be. (Albert Einstein)

Interactive plotting libraries: topic introduction

In this part of the course, we will cover the following concepts:

- Discover different functions to build interactive visualizations
- Visualize data with `highcharter`

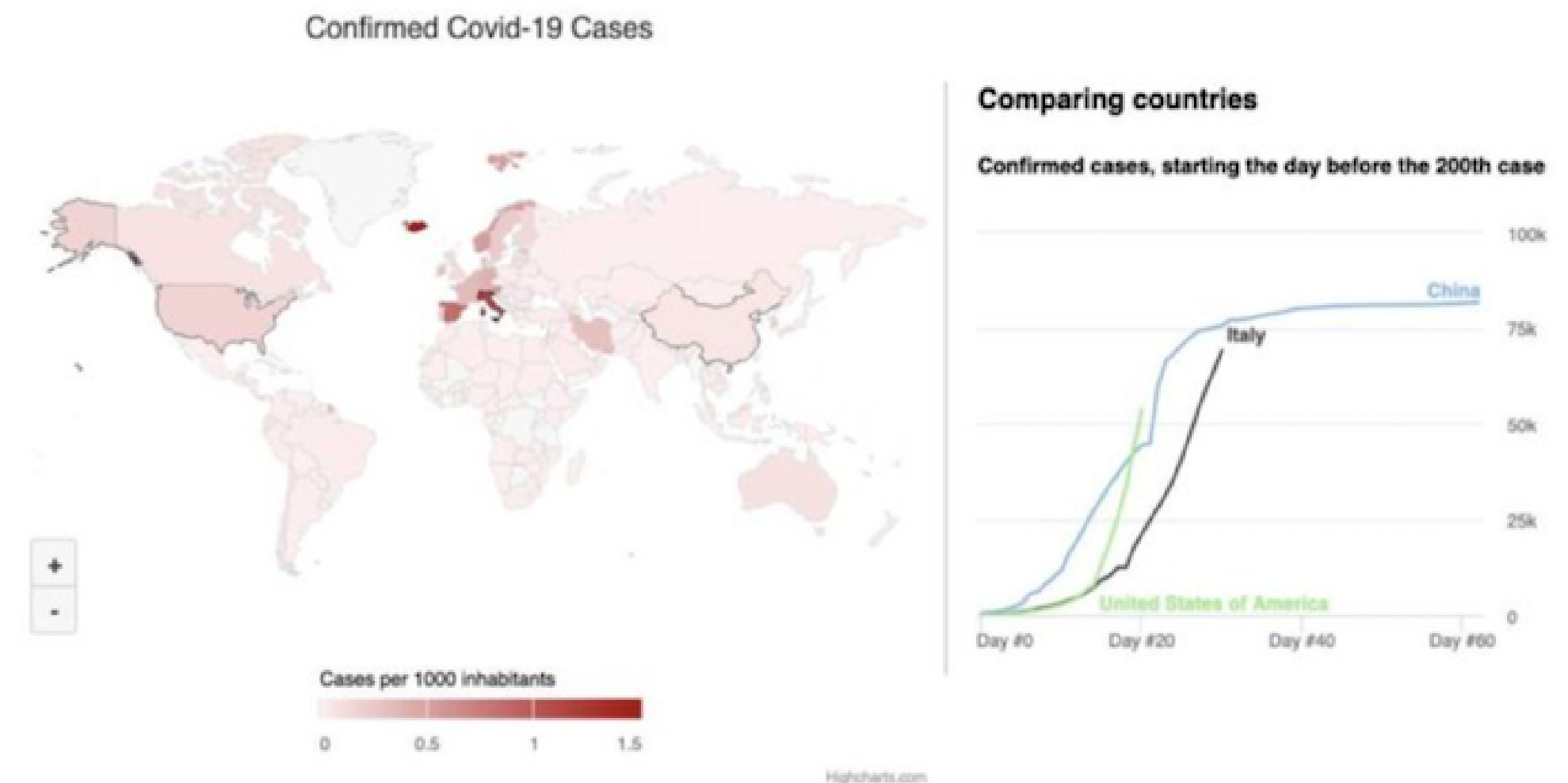
Warm-up

- During the COVID-19 pandemic, the demand for creating quick visualizations to put data into perspective rose quickly
- Before we begin, *read this article about data visualization* and check out an interactive visualization dashboard

A Small Covid-19 Dashboard from Scratch

by [Torstein Honsi](#) and [Vidar Brekke](#)

[Blog Posts](#) [Data Journalism](#) [Data Science](#) [Highcharts](#) [Highmaps](#) [Tutorials](#) [0 comments](#)



Module completion checklist

Objective	Complete
Install the highcharter package and discuss its application to build interactive visualizations	
Create a scatterplot using highcharter with tidy data	

Interactive visualizations with `highcharter`

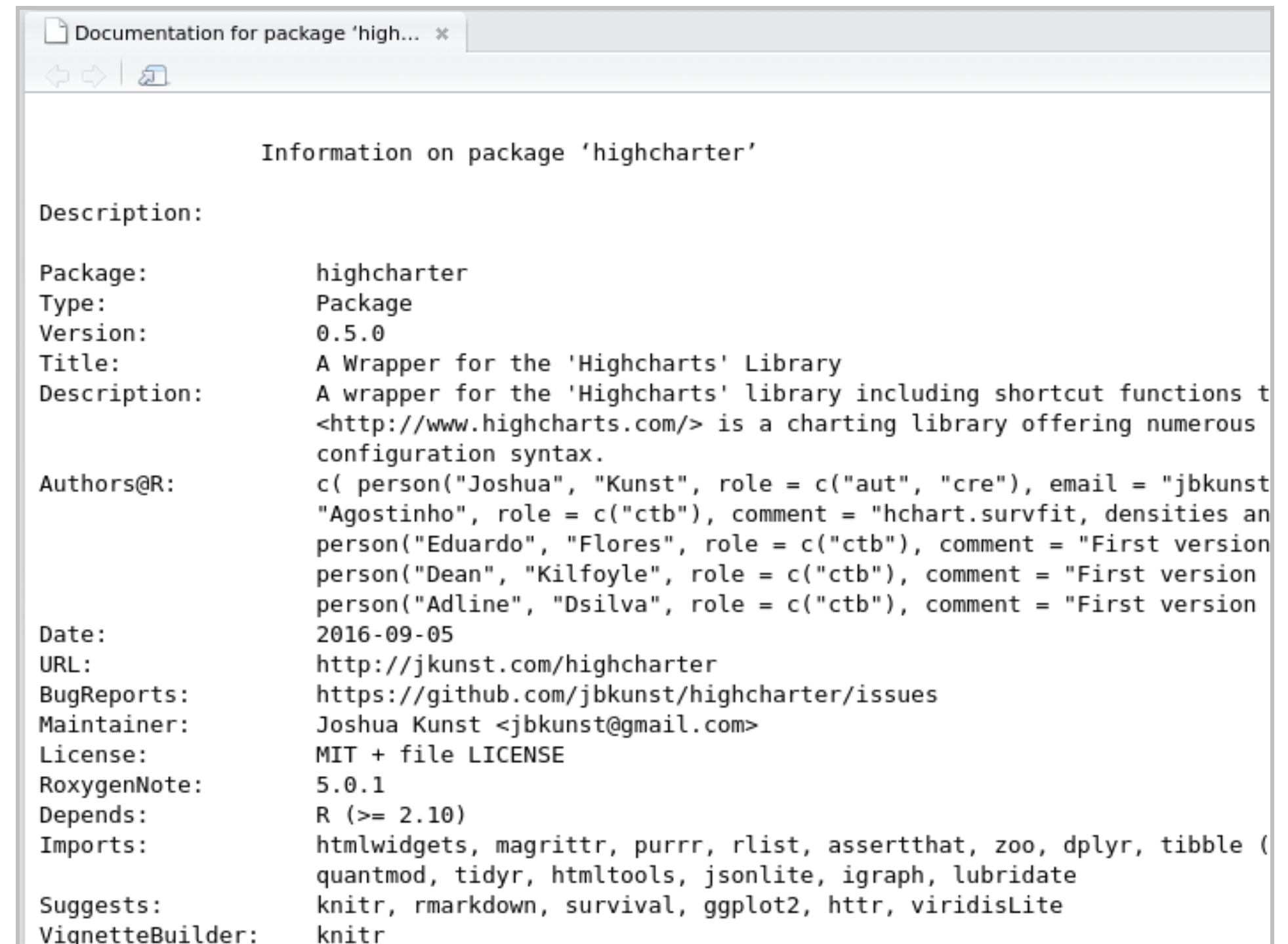
- Highcharter is an R wrapper that allows R users to tap into one of the most comprehensive data visualization JavaScript-based libraries
- Though free for individual research and non-profit purposes, there are some restrictions
- You may need a license to integrate it into a software or organization-wide products
- For more information, *refer to Highcharter's website*

```
> library(highcharter)
Highcharts (www.highcharts.com) is a Highsoft software product which is
not free for commercial and Governmental use
> |
```

Installing highcharter

- Let's install the package and check its documentation

```
# Install `highcharter` package.  
install.packages("highcharter")  
  
# Load the library.  
library(highcharter)  
  
# View documentation.  
library(help = "highcharter")
```



The screenshot shows the R documentation window for the 'highcharter' package. The window title is 'Documentation for package 'high...'. The main content is titled 'Information on package 'highcharter''. It includes a 'Description:' section, a list of package details (Package, Type, Version, Title, Description), 'Authors@R:' information, and various other fields like Date, URL, BugReports, Maintainer, License, RoxygenNote, Depends, Imports, Suggests, and VignetteBuilder.

Information on package 'highcharter'

Description:

Package: highcharter
Type: Package
Version: 0.5.0
Title: A Wrapper for the 'Highcharts' Library
Description: A wrapper for the 'Highcharts' library including shortcut functions to <http://www.highcharts.com/> is a charting library offering numerous configuration syntax.

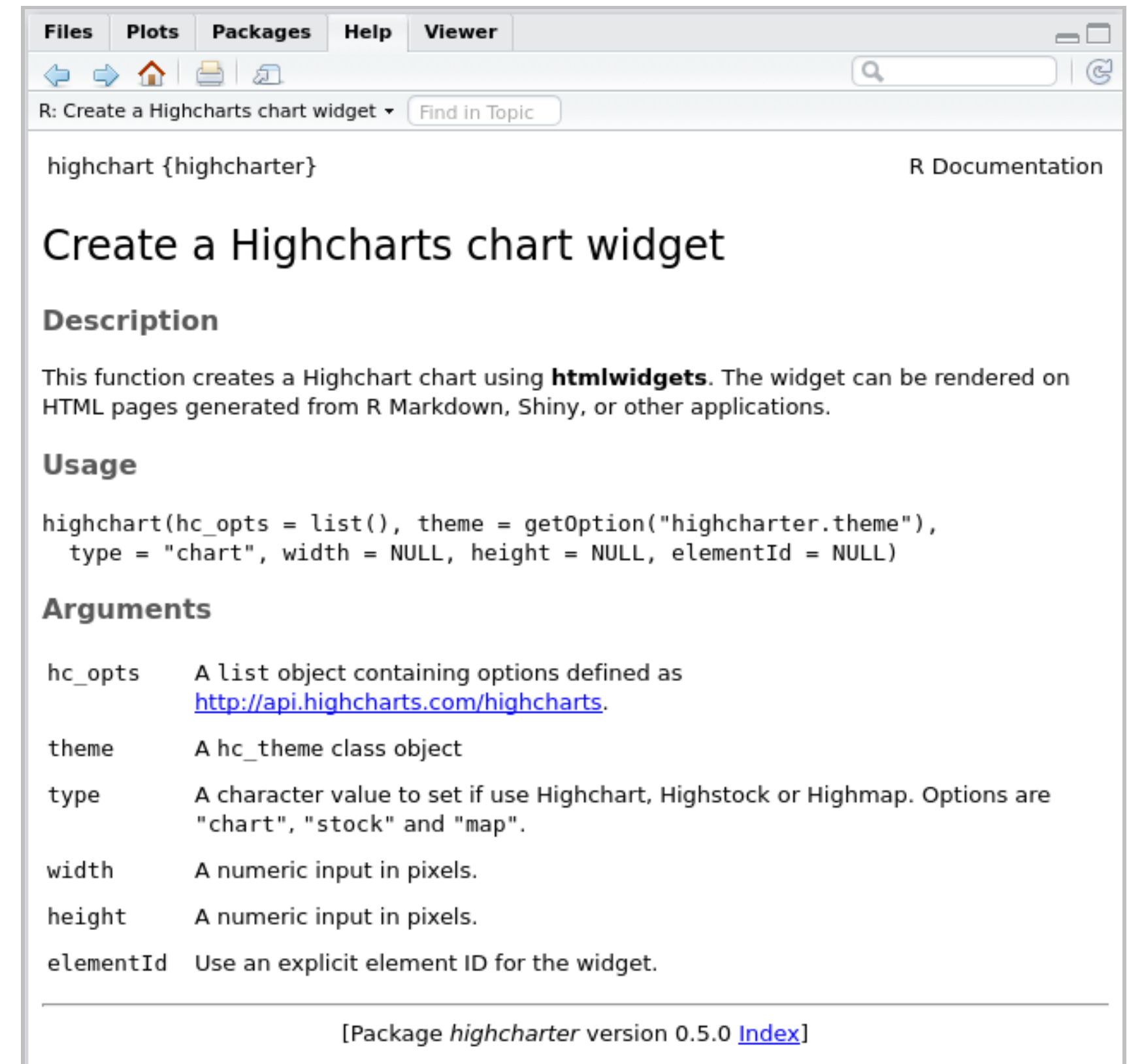
Authors@R: c(person("Joshua", "Kunst", role = c("aut", "cre"), email = "jbkunst@jku.edu", "Agostinho", role = c("ctb"), comment = "hchart.survfit, densities and person("Eduardo", "Flores", role = c("ctb"), comment = "First version person("Dean", "Kilfoyle", role = c("ctb"), comment = "First version person("Adline", "Dsilva", role = c("ctb"), comment = "First version

Date: 2016-09-05
URL: <http://jkunst.com/highcharter>
BugReports: <https://github.com/jbkunst/highcharter/issues>
Maintainer: Joshua Kunst <jbkunst@gmail.com>
License: MIT + file LICENSE
RoxygenNote: 5.0.1
Depends: R (>= 2.10)
Imports: htmlwidgets, magrittr, purrr, rlist, assertthat, zoo, dplyr, tibble (quantmod, tidyr, htmltools, jsonlite, igraph, lubridate
Suggests: knitr, rmarkdown, survival, ggplot2, httr, viridisLite
VignetteBuilder: knitr

Using `highchart()` function

?highchart

- To create a plot, we need to call the main plotting function `highchart()`
- The function doesn't need any required arguments
- The graphic parameters and plotting options can be specified within the layers



The screenshot shows the R Documentation window for the `highchart` function. The window has a menu bar with 'Files', 'Plots', 'Packages', 'Help', and 'Viewer'. Below the menu bar is a toolbar with navigation icons and a search bar. The main content area displays the function signature `highchart {highcharter}` and the title 'Create a Highcharts chart widget'. The 'Description' section states that the function creates a Highchart chart using **htmlwidgets**. The 'Usage' section shows the function call: `highchart(hc_opts = list(), theme = getOption("highcharter.theme"), type = "chart", width = NULL, height = NULL, elementId = NULL)`. The 'Arguments' section lists the parameters: `hc_opts` (A list object containing options defined as <http://api.highcharts.com/highcharts>), `theme` (A `hc_theme` class object), `type` (A character value to set if use Highchart, Highstock or Highmap. Options are "chart", "stock" and "map".), `width` (A numeric input in pixels), `height` (A numeric input in pixels), and `elementId` (Use an explicit element ID for the widget). At the bottom, it indicates the package version: [Package *highcharter* version 0.5.0 [Index](#)].

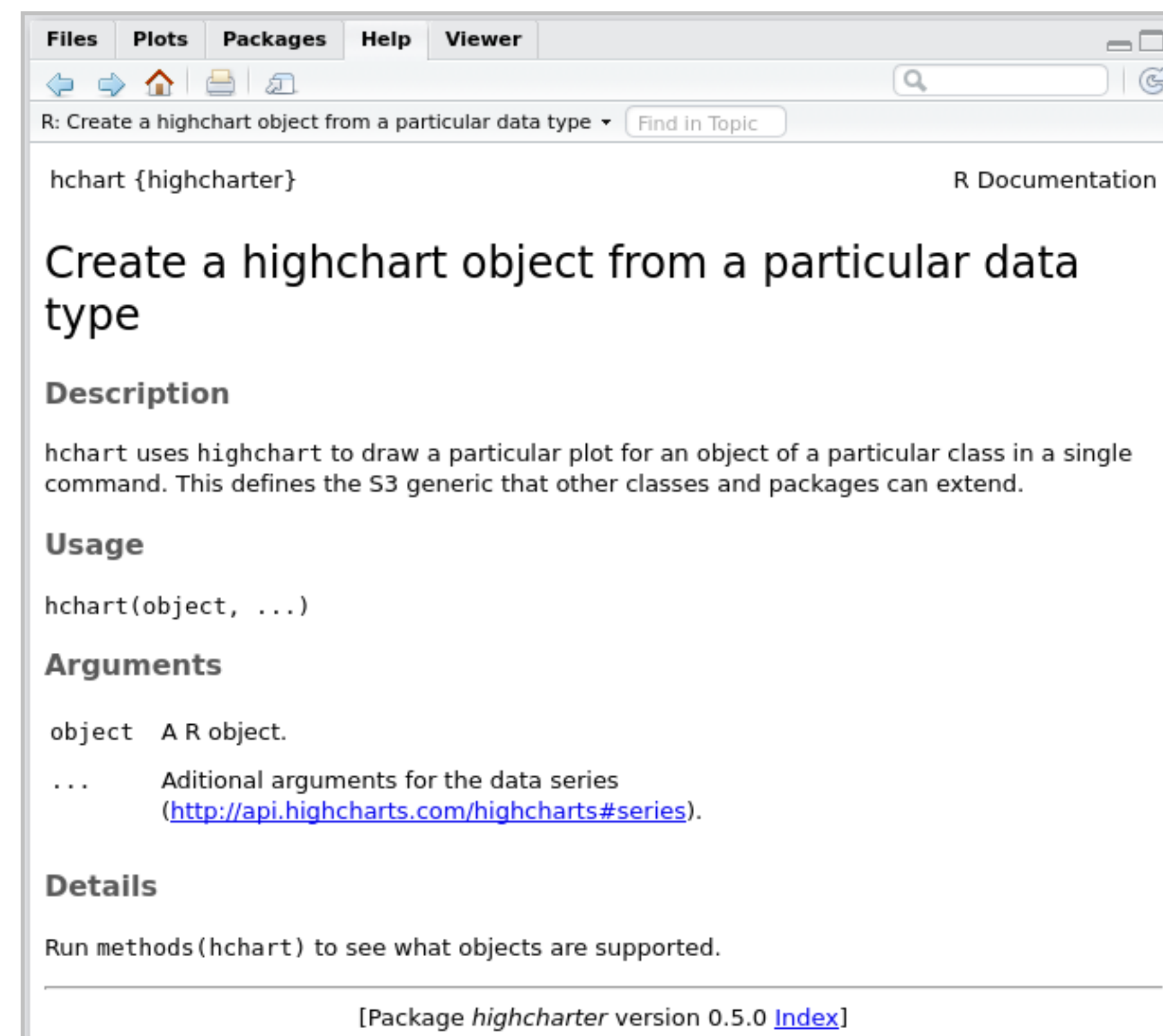
hchart () vs highchart ()

- hchart is a shorthand version of the highchart function

?hchart

```
hchart(Some_data,           #<- dataset to use
        "plot_type",       #<- plot type to use
        hcaes(x = variable1, #<- x-axis mapping
               y = variable2, #<- y-axis mapping
               group = variable3, #<- group by
               ...))
```

- It takes the following arguments:
 - a **dataset** to use
 - the **type of plot** to create (e.g., scatter, bar, column, line, etc.)
 - hcaes (i.e., highchart **aesthetics**) for mapping variables as layers (just as with ggplot2)



The screenshot shows the R Documentation window for the `hchart` function. The window has a menu bar with 'Files', 'Plots', 'Packages', 'Help', and 'Viewer'. Below the menu bar is a search bar and a 'Find in Topic' button. The main content area is titled 'hchart {highcharter}' and 'R Documentation'. The title is followed by the subtitle 'Create a highchart object from a particular data type'. Below this is the 'Description' section, which states that `hchart` uses `highchart` to draw a particular plot for an object of a particular class in a single command. This defines the S3 generic that other classes and packages can extend. The 'Usage' section shows the function signature `hchart(object, ...)`. The 'Arguments' section lists the arguments: `object` (A R object.) and `...` (Additional arguments for the data series (<http://api.highcharts.com/highcharts#series>)). The 'Details' section states 'Run methods(hchart) to see what objects are supported.' At the bottom, it says '[Package highcharter version 0.5.0 [Index](#)]'.

Layers in Highcharter: series

- The highcharter library has its own vocabulary
- Each new data / graphic layer in highcharter is called a **series**
- Series can be of different types; some common ones are listed below:

Highcharter series type	Plot type
scatter	scatterplot
line	line graph
boxplot	boxplot
column	bar plot
bar	horizontal bar plot
histogram	histogram
area	density

Module completion checklist

Objective	Complete
Install the highcharter package and discuss its application to build interactive visualizations	✓
Create a scatterplot using highcharter with tidy data	

Directory settings

- In order to maximize the efficiency of your workflow, use the `box` package and encode your directory structure into `variables`
- Let the `main_dir` be the variable corresponding to your materials folder

```
# Set `main_dir` to the location of your materials folder.  
  
path = box::file()  
main_dir = dirname(dirname(path))
```

Directory settings (cont'd)

- We will store all datasets in the `data` directory inside the `materials` folder in your environment; hence we will save their path to a `data_dir` variable
- We will save all the plots in the `plots` directory corresponding to `plot_dir` variable
- To append one string to another, use `paste0` command and pass the strings you would like to paste together

```
# Make `data_dir` from the `main_dir` and  
# remainder of the path to data directory.  
data_dir = paste0(main_dir, "/data")  
# Make `plots_dir` from the `main_dir` and  
# remainder of the path to plots directory.  
plot_dir = paste0(main_dir, "/plots")
```

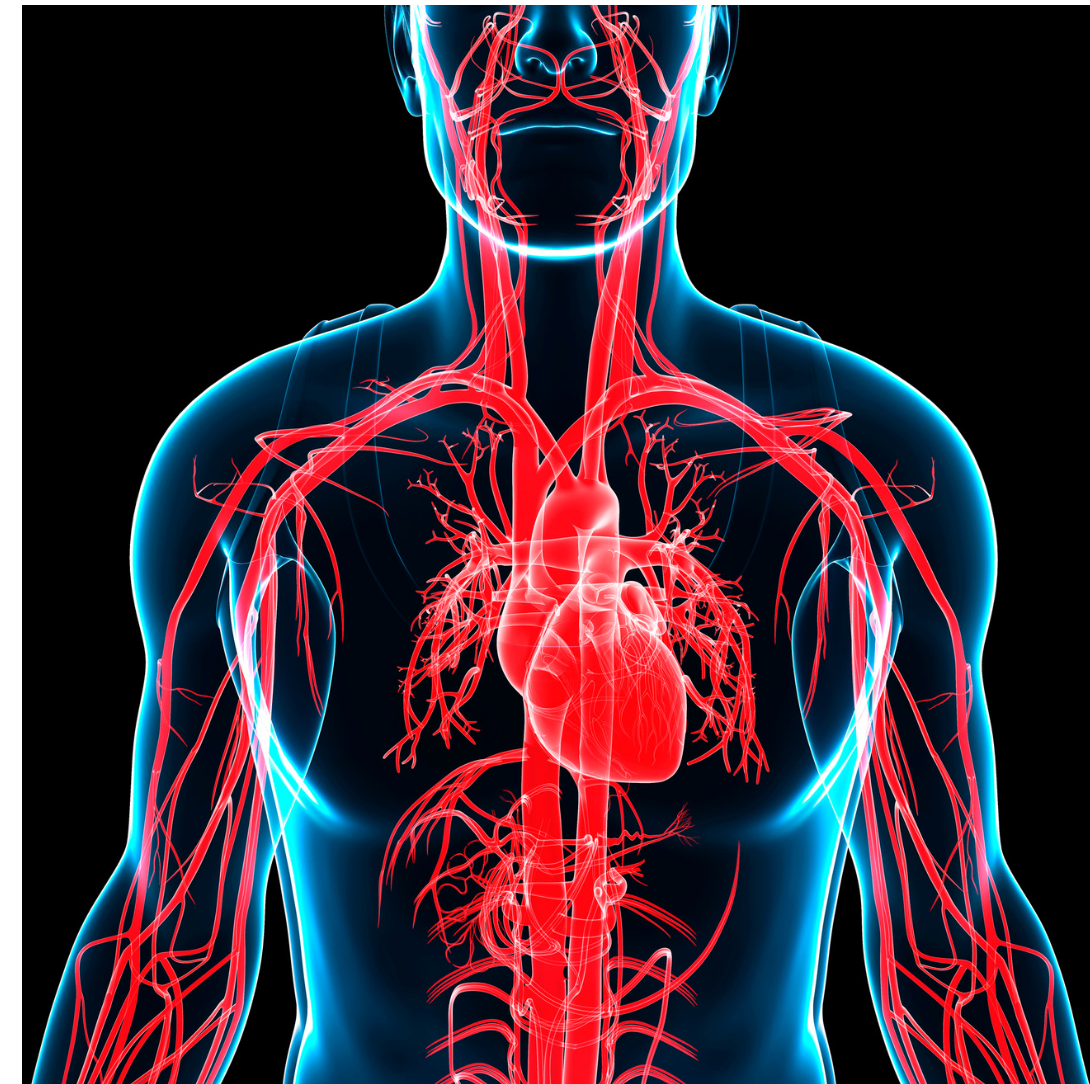
Introducing HDS data set

- We will begin by exploring a dataset called `healthcare-dataset-stroke-data`
- This dataset contains information about age, gender, hypertension, bmi, and other parameters to know the chances of getting a stroke
- The goal is to understand how different variables in the dataset affect the chances of a person suffering from a stroke
- The dataset has 12 characteristics (columns), of which:
 - **10 columns** relate to the **quality and characteristics** of the life of different people
 - The **stroke column** represents whether the people had a stroke or not

Load HDS dataset

- Let's load the HDS dataset from our `data_dir` into R's environment and subset it

```
# Read CSV file called "healthcare-dataset-  
stroke-data.csv"  
HDS = read.csv(file =  
file.path(data_dir, "/healthcare-dataset-stroke-  
data.csv"), #<- provide file path  
              header = TRUE,                #<- if  
file has header set to TRUE  
              stringsAsFactors = FALSE) #<- read  
strings as characters, not as factors
```



Subset data

- In this module, we will explore a dataset subset, including the following variables:
 - age
 - bmi
 - average_glucose_level and
 - stroke

	age	bmi	avg_glucose_level	stroke
1	67	36.60000	228.69	1
2	61	28.89324	202.21	1
3	80	32.50000	105.92	1
4	49	34.40000	171.23	1
5	79	24.00000	174.12	1
6	81	29.00000	186.21	1
7	74	27.40000	70.09	1
8	69	22.80000	94.39	1
9	59	28.89324	76.15	1
10	78	24.20000	58.57	1
11	81	29.70000	80.43	1
12	61	36.80000	120.46	1
13	54	27.30000	104.51	1
14	78	28.89324	219.84	1
15	79	28.20000	214.09	1

Showing 1 to 15 of 5,110 entries, 4 total columns

Prepare Data

- But before sub-setting the data, let's handle the missing data in the dataset
- Then convert `bmi` into a numeric column followed by imputing the missing values with the mean

```
HDS$bmi <- as.numeric(as.character(HDS$bmi)) ##converting bmi column to numeric
# NA imputation
# we can use is.na() function to know about NA values
HDS$bmi[is.na(HDS$bmi)] <- mean(HDS$bmi, na.rm=TRUE) # Replacing na values of bmi column with it's
mean bmi
```

Prepare Data

- Let's tidy the data and transform it from a wide to a long format
- This will be especially useful later for univariate visualizations

```
library(tidyverse)
```

```
# Now Let's make a vector of column indices we would like to save.
```

```
column_ids = select(HDS, age, bmi, avg_glucose_level, stroke)  
HDS_subset = column_ids
```

Create a subset

- Now let's create a different subset to help us build a scatterplot and inspect the head

```
# Prep data for scatterplot

HDS_subset_long = HDS_subset %>%
  gather(-age, #<- gather all variables but `age`
         key = "variable",
         value = "value") %>%
  # All other transformations we've done before.
  group_by(variable) %>%
  mutate(norm_value = value/mean(value, na.rm = TRUE))

head(HDS_subset_long)
```

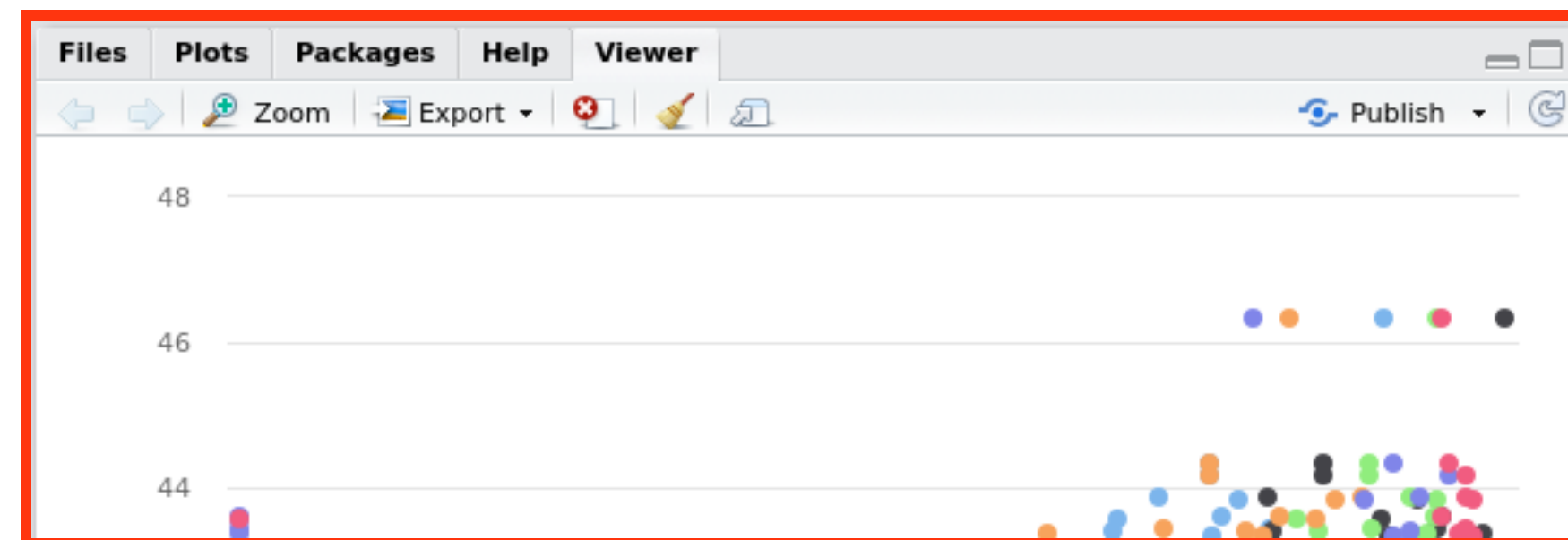
```
# A tibble: 6 x 4
# Groups:   variable [1]
   age variable value norm_value
<dbl> <chr>    <dbl>    <dbl>
1    67 bmi      36.6      1.27
2    61 bmi      28.9       1
3    80 bmi      32.5      1.12
4    49 bmi      34.4      1.19
5    79 bmi       24      0.831
6    81 bmi       29      1.00
```

Construct a scatterplot using `hchart`

- To construct a scatterplot, we use the `hchart()` function and pass the **data**, **plot type** (`scatter`), and **aesthetics** to it as arguments

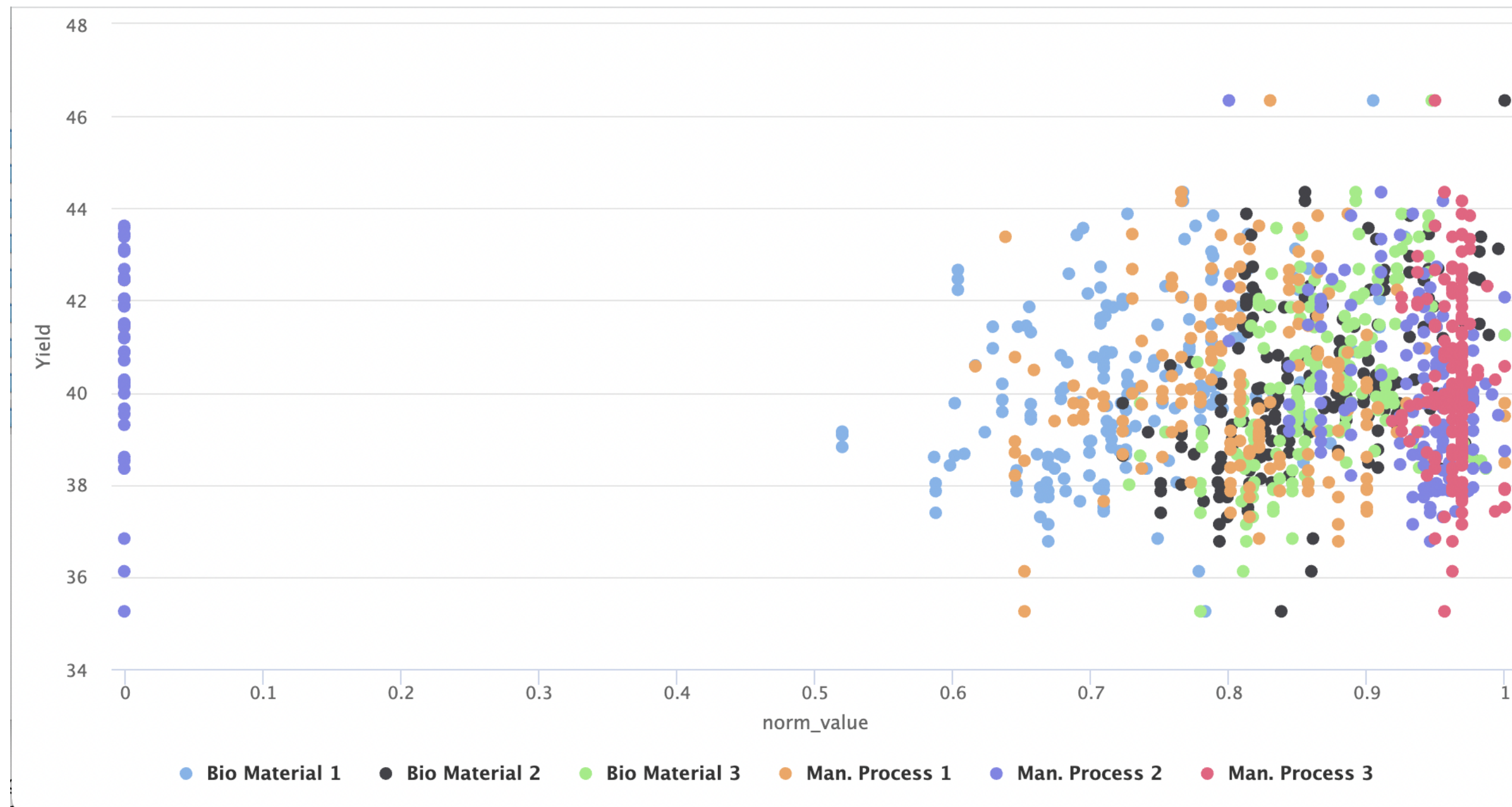
```
# Construct an interactive scatterplot.
scatter_interactive =
  hchart(HDS_subset_long,      #<- name the plot
         "scatter",           #<- set data
         hcaes(x = norm_value, #<- plot type "scatter"
               y = age,         #<- set aesthetics to map x-axis
               group = variable)) #<- set aesthetics to map y-axis
                                   #<- group by
```

- In R, interactive charts appear in the `Viewer` pane, right next to the `Help` tab



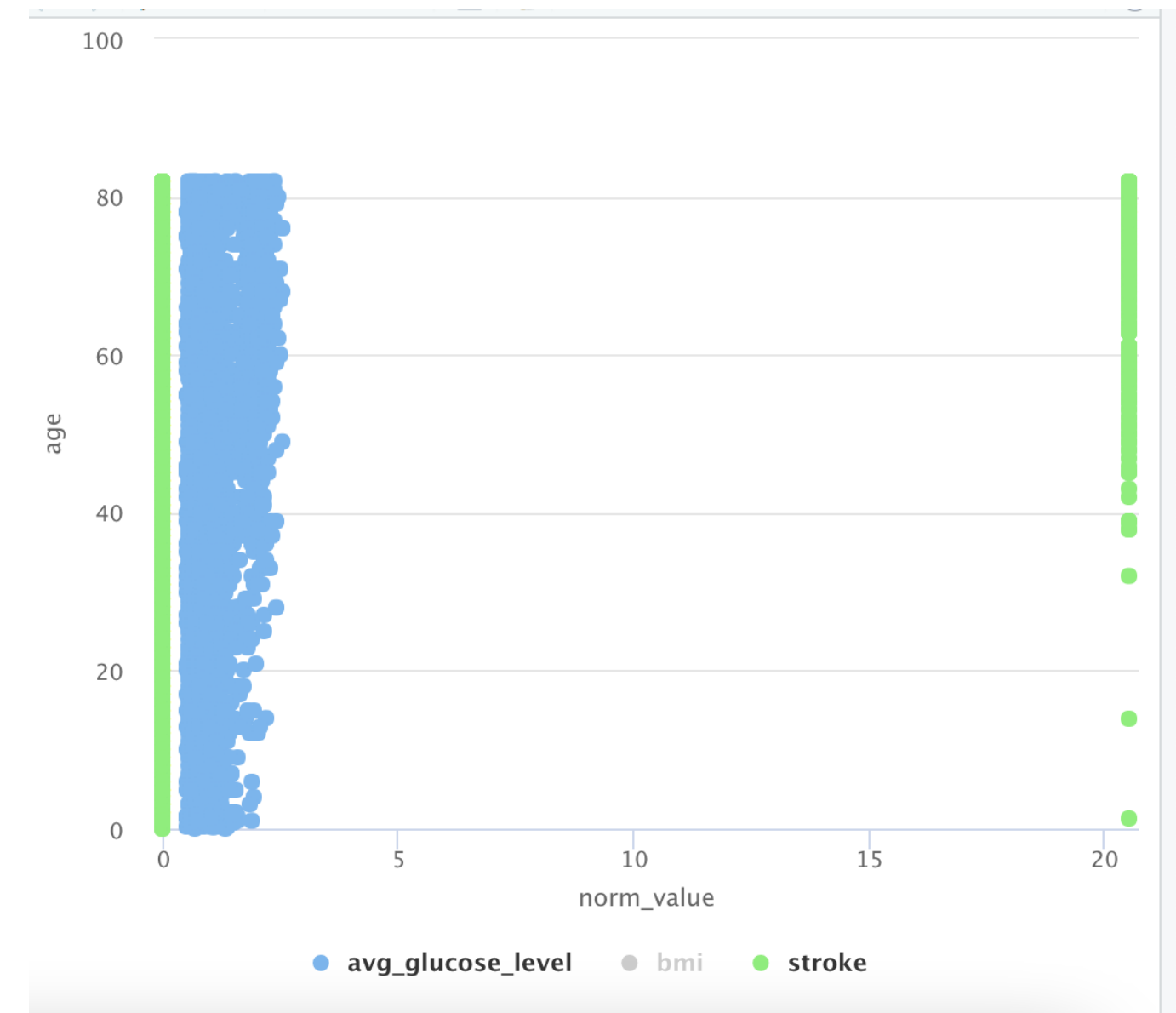
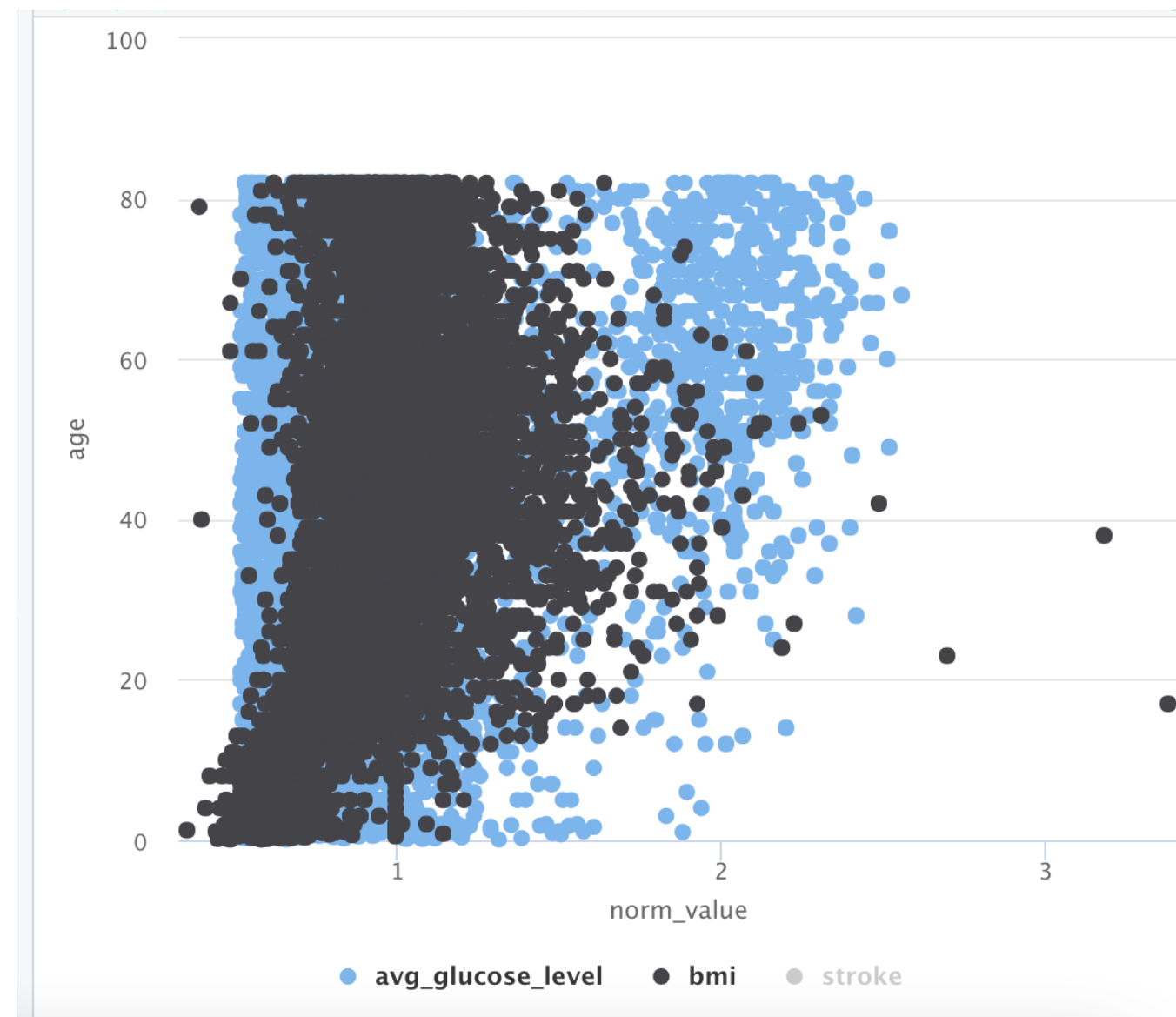
Construct a scatterplot using `hchart` (cont'd)

```
scatter_interactive
```



Selecting categories

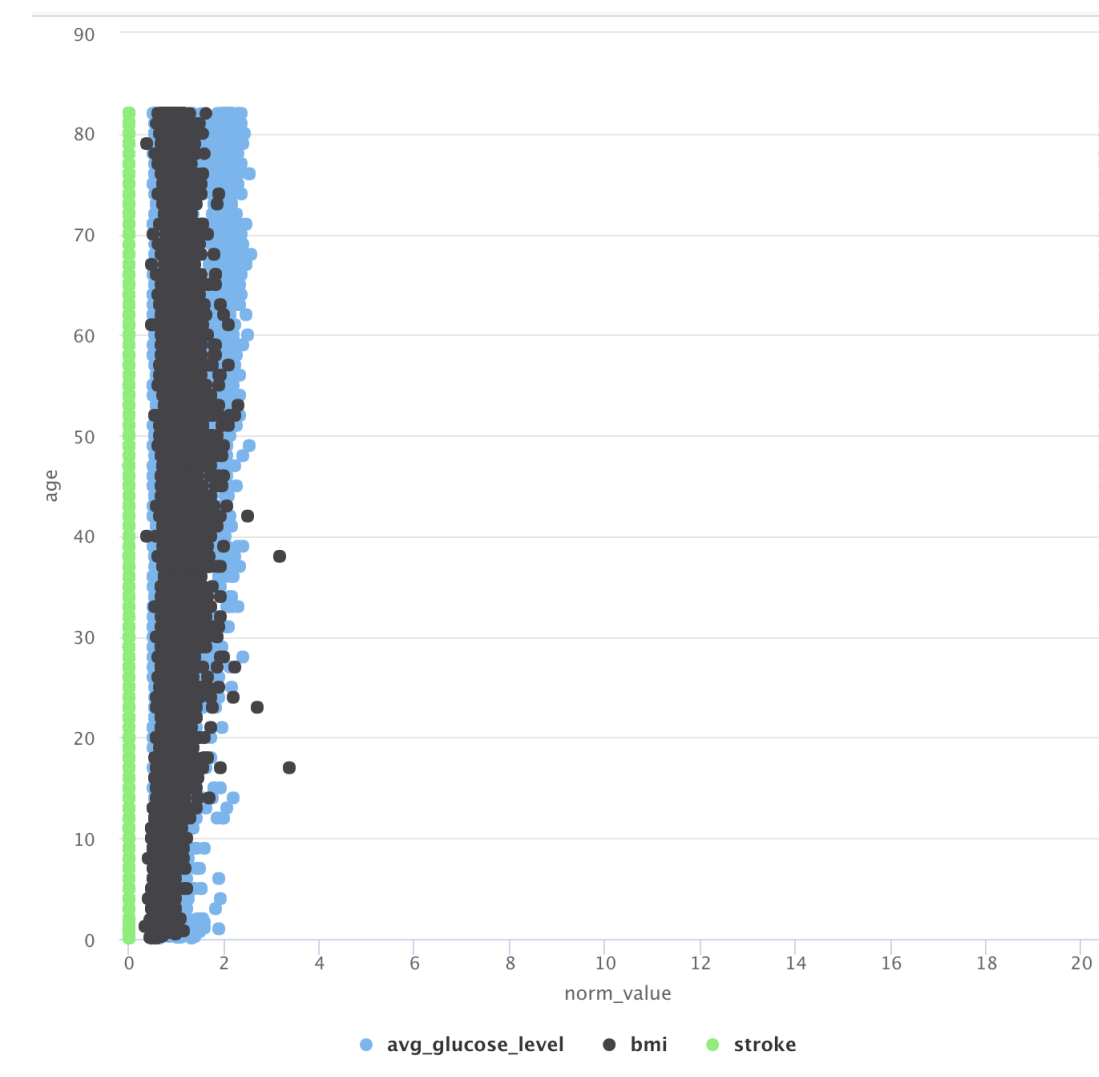
- Every plotted category seen in the legend is a series in highcharter
- When `hchart()` detects more than one category, it auto-colors by series
- We can interactively select and de-select which series to display by clicking on the series names in the legend



Customizing plots with the pipe operator

- You can add a new option or layer using the pipe operator (`%>%`)
- The `hc_chart()` function also controls global chart options like zoom, size, and theme
- Let's zoom in on our plot by passing the `zoomType` argument to `hc_chart()`
 - `xy` zoom allows zooming across both `x` and `y` axes

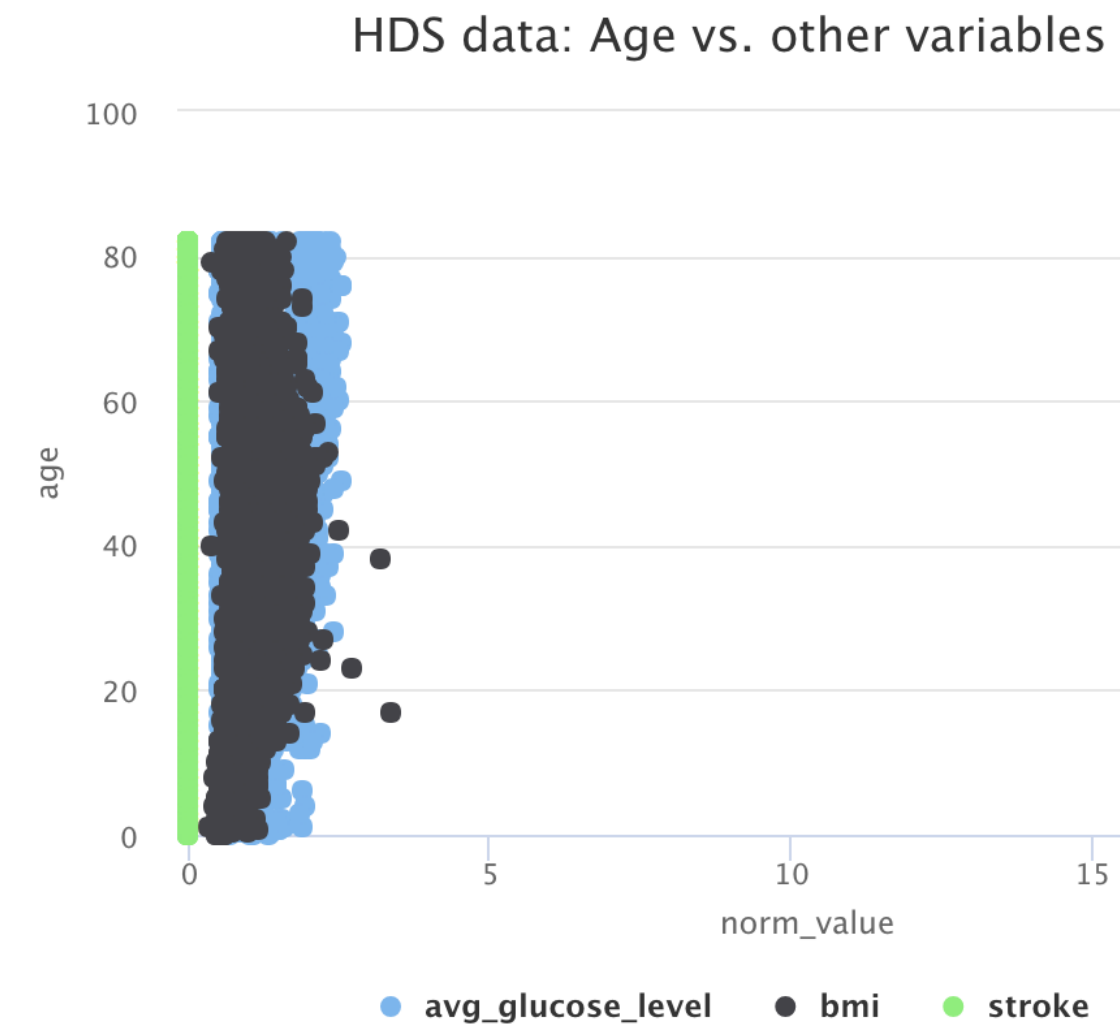
```
# Pipe chart options to original chart.  
scatter_interactive = scatter_interactive %>%  
  # Use chart options to specify zoom.  
  hc_chart(zoomType = "xy")  
  
scatter_interactive
```



Adding a title

- Use the `hc_title()` function to add a title to highcharter plots

```
# Pipe chart options to original chart.  
scatter_interactive = scatter_interactive %>%  
  # Add title to the plot.  
  hc_title(text = "HDS data: Age vs. other  
variables")  
  
scatter_interactive
```



Knowledge check



Exercise



You are now ready to try tasks 1-4 in the Exercise for this topic

Module completion checklist

Objective	Complete
Install the highcharter package and discuss its application to build interactive visualizations	✓
Create a scatterplot using highcharter with tidy data	✓

Interactive plotting libraries: topic summary

In this part of the course, we have covered:

- Discovering different functions to build interactive visualization
- Visualizing data with `highcharter`

Congratulations on completing this module!

