An Interactive Tool to Visualize Results from Uncertainty Quantification

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Abstract

Uncertainty quantification is a class of methods used to simulate the ways that variations to inputs of a system can potentially change the end state of that system. This article outlines the implementation of an interactive tool to visualize the results of an uncertainty quantification analysis. The interactive tool is a deployable web application built with the R Shiny framework. Users can create, adjust, and save custom visualizations to assist in interpreting and presenting results.

Key Words: Interactive Visualization; shiny R Package

1. Introduction

Uncertainty quantification is a methodological framework used with some frequency in engineering analysis (Ewing et al., 2018). It is used to understand how variability within the parameters (i.e. inputs) to some system impact the end state of that system. Engineering analysts use uncertainty quantification to assess and find the balance between design sufficiency and design efficiency. This is particularly important in fields where large-scale prototypes, testing, and data collection are extremely expensive. Engineers in these types of applications are increasingly relying on computational simulations to assess system designs.

In the following article, I outline the implementation of an interactive tool to visualize uncertainty quantification results. The outline of this article will proceed as follows: In Section 2, a brief overview of uncertainty quantification is given. In Section 3, the methods by which the visualization tool was developed are described. In Section 4, features of the resulting web application is described. In Section 5, challenges with and future enhancements to the web application are discussed. The code written to generate the shiny app is included in the Appendix.

2. Uncertainty Quantification Overview

The following description of the uncertainty quantification algorithm is adapted from Ewing et al. (2018). First, a few terms and definitions will be described.

system response quantity (SRQ): A parameter of particular interest directly related to the engineering system in question. The SRQ is the output (i.e. prediction) from the engineering model.

engineering model: A mathematical model that defines the relationship between the parameters (model inputs) and SRQ (model output).

aleatory uncertainty: Uncertainty resulting from randomness inherent to a given parameter. Gaining more knowledge about the parameter will not reduce the uncertainty of the parameter.

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epistemic uncertainty: Uncertainty resulting from a lack of knowledge about a given parameter. Gaining more knowledge about the parameter could reduce the uncertainty of the parameter.

The first step of uncertainty quantification is to identify sources of uncertainty (model parameters) and classify them as either aleatory or epistemic. This classification process has been somewhate debated in literature (Kiureghian and Ditlevsen, 2009), and will not be discussed as it is outside the scope of this project. Once these uncertainties related to the model parameters have been identified and classified as aleatory or epistemic, the uncertainty for each parameter must somehow be described. Traditionally, epistemic uncertainties have been described by an interval over which any value in the interval is equally likely, while aleatory uncertainties have been assigned probability distributions. Some more recent publications (Ewing et al., 2018) propose that all uncertainties, aleatory or epistemic, be described using probability distributions. This debate will not be discussed here as it is, again, outside the scope of this project. Once these distributions and/or intervals have all been assigned, they are carried through the model using Monte Carlo techniques.

Let m denote the number of iterations of an outer for loop, and let n denote the number of iterations in an inner for loop. In the outer for loop, values for the epistemic uncertainty parameters are selected randomly from the intervals/distributions assigned. Upon entering the inner loop, values for the aleatory uncertainty parameters are randomly chosen from the distributions assigned. The values chosen for the parameters in both the outer loop and the inner loop are then used as inputs in the engineering model to calculate a value for the SRQ. This value is stored and the inner loop continues running for the rest of the n-1 iterations. All of the n SRQ values calculated from the n iterations of the inner loop are used to calculate an empirical cumulative distribution function (CDF) of SRQ values and the CDF is stored. The outer loop then begins its second iteration, and new values for the empirical uncertainty parameters are chosen. The inner loop then runs another n iterations, producing another CDF. This process continues until the outer loop has run all of its m iterations.

At this point, there will be m empirical CDFs that have been calculated, representing various possible realized values of the SRQ. These CDFs can then be plotted and interpreted. Ewing et al. (2018) suggests constructing a "P-box". This P-box is found by calculating a lower percentile (e.g. the 5th percentile) and an upper percentile (e.g. the 95th percentile) of the CDF ensemble. This P-box can then be interpreted in several ways, including (1) selecting a SRQ value and extracting a probability interval, and (2) selecting a probability value and extracting an SRQ interval.

3. Methods

In order to assist analysts in visualizing and interpreting results from an uncertainty quantification analysis as described in Section 2, an interactive visualization tool was developed. Since the actual uncertainty quantification analysis can be carried out with greater speed and computational power elsewhere, this implementation does not include the Monte Carlo portion of the algorithm described in Section 2. The web application was built using the shiny R package (Chang et al., 2018) and the code for this application is contained in two files. The first file, ui.R (short for 'user interface') contains the code that controls the layout of the various panels and panes in the application, as well as the placement of the user interface elements (i.e. text inputs, sliders, etc.). The server.R file contains the computational code that performs calculations, data manipulation, and generates the plots. The source code is included in the Appendix.

3.1 R Packages

The following software packages were used in the development of the web application:

- **R:** The code for the web application was written in the R programming language (R Core Team, 2018).
- **RStudio:** Rstudio (RStudio Team, 2016) is an integrated development environment that was used to both develop and deploy the web application.
- shiny: The shiny package and framework (Chang et al., 2018) constitues the backbone of this project. shiny provides a way to create and deploy web applications through RStudio (RStudio Team, 2016). It also contains the implementations for all user-interface components (toggle buttons, check boxes, numeric inputs, sliders, etc.). The shinydashboard package (Chang and Borges Ribeiro, 2018) was also used as an aesthetic wrapper around the shiny framework. The shiny package was be used to create and will be used to deploy the web application.
- **ggplot2:** The plotting functionality of the ggplot2 package (Wickham, 2016) was used to generate the actual visualization and to add, remove, or adjust components on the plot.
- **shinydashboard:** The shinydashboard package (Chang and Borges Ribeiro, 2018) was used as an aesthetic wrapper to the web application to improve its professional appearance.
- **shinyalert:** The shinyalert package (Attali and Edwards, 2018) was used to create a pop-up message when the user takes certain actions.
- **shinyWidgets:** The shinyWidgets package (Perrier et al., 2019) was used to create custom toggle buttons.

Since the existing R packages already implemented most of the tools needed to generate the plot and implement user interface elements, the primary task on this project was to seamlessly combine elements from the packages above (particularly the shiny, ggplot2 packages) to create a user-friendly interactive visualization tool.

4. Results

The resulting web application currently several main features implemented. The web application is currently deployed at https://misaac.shinyapps.io/UQViz/. The following subsections outlines the basic functionality.

4.1 Data Source

The 'Data Source' tab contains the controls that allow the user to select either a sample data set (provided for convenience) or to upload their own .csv file. If data from a file is to be used, the x-values over which the CDFs are evaluated should be the first column, and the CDF y-values in the following columns.

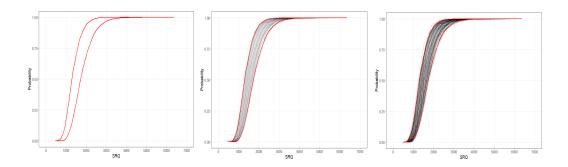


Figure 1: Visualization with P-box shown and CDfs hidden (left); Visualization with P-box and CDFs both shown with high transparency (center) and low transparency (right)



Figure 2: Screenshot from the web application of visualization option control panel

4.2 Visualization Options

Several visualization options are provided to allow customizable graphics to be created. The 'Show/Hide P-box' and the 'Show/Hide CDFs' buttons toggle back and forth between showing and hiding the red P-box and the black ensemble of CDFs. Note that showing the CDF ensemble may (depending on the size of the uploaded file) significantly increase plot rendering times.

The P-box lower and upper percentiles default to 0.05 and 0.95. Users can modify this by inputting the desired lower and upper percentile to be used in the visualization.

Moving the 'CDF Transparency' slider will make each individual CDF trace more or less transparent. A slider value of 1 corresponds to completely opaque, while a slider value of 0 corresponds to completely transparent. This mitigates the issue of overplotting that can often arise when hundreds or thousands traces are shown on the same plot. Allowing for some transparency allows the user to visualize the density of regions within the CDF ensemble. Figure 1 shows plots with several possible visualization options selected. A screenshot of the visualization option control panel is shown in Figure 2.

To aid with interpretation, both probability intervals and SRQ intervals can be extracted from the P-box shown in the visualization, as described in Ewing et al. (2018). To extract a probability interval, the user selects an SRQ value at which a line is drawn vertically. The intersections of the vertical line with the P-box create the lower and upper endpoints for the probability interval. In a similar process, the user can also extract a SRQ interval by selecting a probability value at which a line is drawn horizontally. The intersections of the horizontal line with the P-box create the SRQ interval. This process is illustrated in Figure 3.

A title and can be added to the plot and the x-axis label can be changed from the default label of 'SRQ'. This allows users to futher customize the visualization to a specific domain or context.

Users can download and save a .png version of the current rendering of the visualiza-

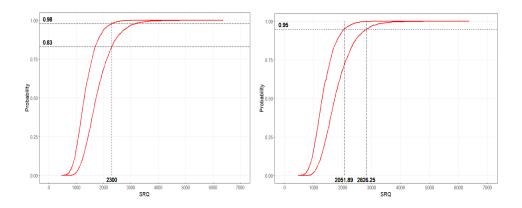


Figure 3: Extracting a probability interval (left); Extracting an SRQ interval (right)

tion by providing a file name and clicking the 'Download' button.

5. Discussion

5.1 Challenges

One main challenge with this visualization tool is the time that it would takes to render the plot when the CDF ensemble is shown. The nature of uncertainty quantificataion and this kind of visualization leads to there being hundreds or even thousands of traces on a single plot. In addition, the way the interactive features are implemented through the shiny package causes the plot to completely re-render when user inputs change. Thus, whenever the user selects a new plotting option and the CDFs are currently shown, he or she must wait several seconds (depending on the size of the data set being used) for the plot to re-render. This leads to a less 'interactive' feel for the user. No immediate or direct solution has been found for this issue yet. However, an effective workaround, as described the the User Guide tab on the web application, has been suggested. If the user wishes to display the CDF ensemble on the final version of the plot, it is suggested to adjust all visualization options and controls while the CDFs are hidden, keeping the plot rendering times to a reasonable length. Then, as the last step, show the CDF ensemble. This way, the user will only have to wait once for the long rendering time.

5.2 Future Work

Further work in developing this visualization tool may involve researching other methods for visualizing uncertainty, such as the methods described in Jackson (2008), and implementing several methods for visualization. This would allow users to choose from various visualization methods within the web application to create a graphic that is best suited to a given context. Another extension of this visualization tool could be to use it in a self contained UQ tool, where users could perform the entire UQ algorithm from start to finish. For example, users could define the engineering model to be used, define parameters as aleatory or epistemic, carry out the UQ algorithm, and visualize the results. However, as the web application now stands, it can be used to assist engineers and analysts in visualizing, understanding, and interpreting results from uncertainty quantification analyses.

References

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- Wickham, H., 2016. ggplot2: Elegant Graphics for Data Analysis. Springer-Verlag New York, (http://ggplot2.org).

A. Appendix

A.1 Source Code: ui.R

```
library(shiny)
library (shinydashboard)
library (plotly)
library(shinyalert)
library(shinyWidgets)
cranURL <- "https://cran.r-project.org/web/packages/"</pre>
dashboardPage(
  dashboardHeader(
   title = "UQViz"
  dashboardSidebar(
  dashboardBody(
   useShinyalert(),
    tabsetPanel(
      tabPanel ("Data Source",
               fluidRow(
                 column (
                   width = 3,
                   choices = c("Sample Data",
                                            "File Upload"),
                                selected = "Sample Data"),
                   uiOutput (outputId = "fileInput")
      tabPanel("Visualization Options",
               fluidRow(
                 column (
                   width = 2,
                   offset = 0,
                   verticalLayout(
                    br(),
                     uiOutput (outputId = "pbxBttn"),
                     uiOutput (outputId = "cdfBttn")
                 ),
                 column (
                   width = 2,
                   offset = 0,
                   numericInput(inputId = "pboxLower",
                                label = "Lower P-box Percentile",
                                value = 0.05,
width = 250),
                   numericInput(inputId = "pboxUpper",
                                label = "Upper P-box Perentile.",
                                value = 0.95,
                                width = 250)
                 column (
                   width = 3,
                   offset = 0,
                   sliderInput(inputId = "sliderAlpha",
                               label = "CDF Transparency",
                               value = 0.2,
                               min = 0,
                               max = 1,
                               step = 0.1,
```

```
ticks = FALSE)
           ),
           column (
             width = 1,
             offset = 0,
             checkboxInput(inputId = "extractInt",
                           label = "Extract Interval",
                           value = FALSE)
           ),
           column (
             width = 2,
             uiOutput (outputId = "extractTypeInput"),
             uiOutput (outputId = "valInput")
tabPanel ("Labeling",
         fluidRow(
           column (
             width = 2,
             textInput(inputId = "titleInput",
                       label = "Title",
                       placeholder = "Plot Title",
           column (
             width = 2,
             textInput(inputId = "xaxisInput",
                       label = "X axis Label",
                       placeholder = "X-axis",
                       value = "SRQ") #,
             # textInput(inputId = "yaxisInput",
                         label = "Y axis Label",
                         placeholder = "Y-axis",
                         value = "Probability")
         )),
tabPanel("User Guide",
         h3("Overview"),
         h5("UQViz is an interactive tool that can be used
            to create meaningful visualizations of
            '2D Uncertainty Quantification'
            (for more information on 2D UQ, see ",
            tags$<mark>a</mark>(
              href = paste0("http://verification.asmedigitalcollection",
                             ".asme.org",
                             "/article.aspx?articleid=",
                             "2703289&resultClick=1"),
                   "this article",
                   target = "_blank"),
            "). Users can select data source and adjust various
            visual components and labels. The title and axis
            label can also be customized, and the visualization
            can be downloaded and saved."),
         h3("Data Source"),
         tags$p("The 'Data Source' tab contains controls for
                the user to select the source from
                which data for the visualization will be obtained.
                If ",
                tags$b('Sample Data'),
                "is selected, a small built-in data set is used as
                the data source for the
                visualization. This feature is provided for convenience,
                and can be used for experimentation and
                exploration. If ",
                tags$b('File Upload'), "
                is selected, users can upload a ",
                tags$code(".csv"),
                "file containing
```

```
data to be used in the visualization. This ",
       tags$code(".csv"),
       "file should be formatted such that the 'x' values over
       which the CDFs are evaluated
       are in the first column, and the CDFs are
       in the following columns."
h3("Visualization Options"),
tags$p("Several visualization options are provided to
       allow customizable graphics to be
       created. The ",
       tags$b("Show/Hide P-box"), " and the ",
       tags$b("Show/Hide CDFs"), " buttons toggle
       back and forth between showing and
       hiding the red P-box and the black ensemble of CDFs.
       Note that showing the CDF
       ensemble may (depending on the size of the uploaded
       file) significantly increase
       plot rendering times. If the user wishes to
       display the CDFs on the final version
       of the plot, it is suggested to adjust all
       visualization options and controls
       while the CDFs are hidden, keeping the plot
       rendering times to a reasonable
       length. Then, as the last step, show the CDF
       ensemble. This way, the user will only
       have to wait once for the lengthened rendering time."),
tags$p("The ",
       tags$b("Lower P-box Percentile"),
       " and ",
       tags$b("Upper P-box Percentile"),
       "values can also be adjusted. This will
       shift the boundaries of the P-box on
       the visualization."),
tags$p("Adjusting the ",
       tags$b("CDF Transparency"),
       " slider will adjust the
       transparency of each CDF in the CDF ensemble.
       A value of 0 corresponds to
       completely transparent, while a value
       of 1 corresponds to completely opaque.
       This control will only affect the visualization
       when the CDFs are being shown.
       This feature can be useful when there is a
       high degree of overplotting due to
       a large number of CDFs. Allowing transparency
       on the CDFs can assist in
       visualizing high-density regions on the plot."),
tags$p("When the ",
       tags$b("Extract Interval"),
       " box is checked,
       several more controls appear.
       Extracting an interval is one way that
       P-boxes can be interpreted in practice.
       Users can decide whether to extract an ",
       tags$b("SRQ Interval"),
       " or a ",
       tags$b("Probability Interval"),
       ". If extracting an SRQ interval, users
       will need to select a probablity value; if
       extracting a probability interval, users
       will need to select an SRQ value.
       Labels and dotted lines will be displayed
       on the visualization, showing the
       probability or SRQ value selected and the
       corresponding interval."),
h3("Labeling"),
tags$p("Users also have the capability to choose
       a custom ",
       tags$b("Title"),
       " and a custom ",
```

```
tags$b("X axis Label"),
                   "to futher customize the plot to a
                   specific context or application. Since this
                   visualization was designed to always show
                   probability on the y axis, the y axis label
                   cannot be changed."),
            h3("Saving a Plot"),
            tags$p("Users can enter a ",
                   tags$b("File Name"),
                   " and can click the "
                   tags$b("Download"),
                   " button to download and save a ",
                   tags$code(".png"),
                   " file of the current version of the visualization.
                   These controls are located below the plot.
                   Note that the chosen
                   file name should omit the ",
                   tags$code(".png"),
                   " file extension.")
   tabPanel("References",
            tags$h4("The following software packages
                    were used in development: "),
            tags$div(tags$ul(
              tags$li(tags$span(tags$a(href = "https://www.R-project.org/",
                                       target = "_blank",
                                       "R"))),
              tags$li(tags$span(tags$a(href = "http://www.rstudio.com/",
                                       target = "_blank",
                                        "RStudio"))),
              tags$li(tags$span(tags$a(
               href = paste0(cranURL, "shiny/index.html"),
                target = "_blank",
                "shiny"))),
              tags$li(tags$span(tags$a(
                href = paste0 (cranURL, "ggplot2/index.html"),
                target= "_blank",
                "ggplot2"))),
              tags$li(tags$span(tags$a(
                href = paste0(cranURL, "shinyWidgets/index.html"),
                target = "_blank",
                "shinyWidgets"))),
              tags$li(tags$span(tags$a(
                href = paste0(cranURL, "shinydashboard/index.html"),
                target = "_blank",
                "shinydashboard"))))
  )
 fluidRow(
  box (
     width = 12,
    height = 625,
    plotOutput("uqPlot",
                height = 475),
     fluidRow(
       column (
        width = 3,
         textInput(inputId = "pngName",
                   value = "",
                   placeholder = "file name",
                   label = "File Name (omit file extension)",
                   width = 200),
         downloadButton(outputId = 'savePng')
    )
  )
)
```

A.2 Source Code: server.R

```
# Load needed packages
library (shiny)
library (ggplot2)
library(plotly)
library(shinyalert)
library(shinyWidgets)
## Increase max file upload size
options(shiny.maxRequestSize = 15*1024^2)
# Function to calculate p-box lower and upper bounds
calc_pbox <- function(data, p_upper, p_lower){</pre>
 pbox <- apply(values$data,</pre>
             FUN = quantile,
             probs = c(p_lower, p_upper))
 pbox <- t(pbox)</pre>
 values$pbox <- data.frame(x = values$data[,1],</pre>
                      lower = pbox[,1],
                       upper = pbox[,2])
 return (pbox)
# function to calculate the SRQ value for a given probability level
calc_srg <- function(prob_in){</pre>
 srq_1 <- values$pbox[which.min(abs(values$pbox$lower - prob_in)),]$x</pre>
 srq_u <- values$pbox[which.min(abs(values$pbox$upper - prob_in)),]$x</pre>
 return(c(srq_l, srq_u))
}
# function to calculate the probability value for a probability level
calc_probs <- function(srq_in){</pre>
 prob_l <- values$pbox[which.min(abs(values$pbox$x - srq_in)),]$lower</pre>
 prob_u <- values$pbox[which.min(abs(values$pbox$x - srq_in)),]$upper</pre>
 return(c(prob_l, prob_u))
# Reactive Values#################################
values <- reactiveValues()</pre>
# Read in Sample Data ########################
cdf_arr <- read.csv("data/cdfs.csv", header = FALSE)
cdf_arr <- t(cdf_arr)
cdf_df <- data.frame(cdf_arr)</pre>
values$valmax <- max(cdf_df[,1])</pre>
values$valmin <- min(cdf_df[,1])</pre>
values$userin <- NULL
values$showcdf <- FALSE
values$showpbx <- TRUE</pre>
shinyServer(function(input, output) {
```

```
# toggle button to show/hide cdf ensemble
observeEvent (input$cdfToggle, {
  if (values$showcdf == FALSE) {
    shinyalert(
      title = "",
      callbackR = function(x){
        values$showcdf <- x
      text = paste("Showing the CDF ensemble on the plot may significantly",
                    "increase plot render times. Do you want to continue?"),
      showCancelButton = TRUE,
      confirmButtonText = "Show CDFs",
      cancelButtonText = "Don't Show CDFs",
      confirmButtonCol = "#52D755",
      type = "info"
  } else {
  values$showcdf <- FALSE
})
observeEvent (input$pbxToggle,
  if (values$showpbx == FALSE) {
    values$showpbx <- TRUE</pre>
  } else {
    values$showpbx <- FALSE
})
# toggle button to show/hide p-box
output$pbxBttn <- renderUI({
  if (values$showpbx == TRUE) {
    lbl = "Hide P-box"
    sty = "bordered"
  } else {
  lb1 = "Show P-box"
    sty = "simple"
  actionBttn(inputId = "pbxToggle",
              label = lbl,
              style = sty,
              size = "sm",
              color = "primary")
})
output$cdfBttn <- renderUI({
  if (values$showcdf == TRUE) {
    lbl = "Hide CDFs"
    sty = "bordered"
  } else {
  lbl = "Show CDFs"
    sty = "simple"
  actionBttn(inputId = "cdfToggle",
             label = lbl,
              style = sty,
              size = "sm",
              color = "primary")
})
output$uqPlot <- renderPlot({
  # Use selected data source
  if (input$radioDataSource == "Sample Data") {
    values$data <- cdf_df</pre>
  } else {
    req(input$fileIn)
    df <- read.csv(input$fileIn$datapath,</pre>
                    header = FALSE)
    df <- data.frame(df)</pre>
```

```
values$data <- df
# set up initial plot
req(values$data)
plt <- ggplot (data = values$data,
               aes(values$data[,1]))
# plot cdf ensemble if user selects
if (values$showcdf == TRUE) {
  for(i in names(values$data)[-1]){
    plt <- plt +
      geom_line(aes_string(y = i,
                            color = shQuote("CDFs")),
                 alpha = input$sliderAlpha)
 plt <- plt
# get nice breaks for x-axis
px <- pretty(values$data[,1])</pre>
# Labels and additional plot formatting
plt <- plt +
 xlab(input$xaxisInput) +
  # ylab(input$yaxisInput) +
  ylab("Probability") +
  scale_x_continuous(breaks = px,
                      limits = range(px)) +
  ggtitle(input$titleInput) +
  theme_bw() +
  theme(plot.title = element_text(hjust = 0.5,
                                    size = 18,
                                    face = "bold"),
        panel.grid.minor = element_blank(),
        axis.text = element_text(size = 12),
        axis.title = element_text(size = 14))
# show p-box if user selects
if (values$showpbx == TRUE) {
  pbox <- calc_pbox (values$data,
                     (1 - input$pboxLower),
                      (1 - input$pboxUpper))
  plt <- plt +
    geom\_line(aes(y = pbox[,1], col = "Pbox"), lwd = 1) +
    geom_line(aes(y = pbox[,2], col = "Pbox"), lwd = 1) +
scale_colour_manual(name="", values=c("Pbox"="red"))
# Calculate and show extracted interval on plot
# if user selects
if (input$extractInt == TRUE) {
  req(input$valIn)
  # if interval is an SRQ interval
  if (input$extractType == "SRQ Interval"){
    if (input$valIn < 0){</pre>
      values$userin <- 0
    } else if (input$valIn > 1) {
      values$userin <- 1</pre>
    } else {
      values$userin <- input$valIn</pre>
    srq_vec <- calc_srq(values$userin)</pre>
    plt <- plt +
      geom_hline(yintercept = values$userin, lty = 2) +
      geom_segment (mapping = aes(x = srq_vec[1],
                                   y = 0,
                                   xend = srq_vec[1],
                                   yend = 1),
                    lty = "longdash") +
```

```
geom_segment (mapping = aes (x = srq_vec[2],
                                   y = 0,
                                   xend = srq_vec[2],
                                   yend = 1),
                    lty = "longdash") +
      geom_text(aes(srq_vec[1],0.05,
                     label = round(srq_vec[1],2),
vjust = 3,
                     hjust = 0.5),
                 size = 5) +
      geom_text(aes(srq_vec[2],0.05,
                     label = round(srq_vec[2],2),
vjust = 3,
                     hjust = 0.5),
                 size = 5) +
      geom_text(aes(min(values$data[,1]),
                     values$userin,
                     label = round(values$userin,2),
                     vjust = -0.5,
                     hjust = 2),
                size = 5)
  } else {
    # if interval is a probability interval
    req(input$valIn)
    if (input$valIn < values$valmin) {</pre>
      values$userin <- values$valmin</pre>
    } else if (input$valIn > values$valmax) {
      values$userin <- values$valmax</pre>
    } else {
      values$userin <- input$valIn</pre>
    prob_vec <- calc_probs(values$userin)</pre>
    plt <- plt +
      geom_segment (mapping = aes (x = values$userin,
                                  y = 0,
                                   xend = values$userin,
                                   yend = 1),
                    ltv = 2) +
      geom_hline(yintercept = prob_vec[1],
                  lty = "longdash") +
      geom_hline(yintercept = prob_vec[2],
                  lty = "longdash") +
      geom_text (aes (min (values$data[,1]),
                     prob_vec[1],
                     label = round(prob_vec[1],2),
                     vjust = -0.5,
                    hjust = 2),
                size = 5) +
      geom_text(aes(min(values$data[,1]),
                     prob_vec[2],
                     label = round(prob_vec[2],2),
                     vjust = -0.5,
                    hjust = 2),
                size = 5) +
      geom_text (aes (values$userin,
                     0.05,
                     label = round(values$userin,2),
                     vjust = 3,
                     hjust = 0.5),
                 size = 5)
if (values$showpbx && values$showcdf){
 plt <- plt +
    scale_color_manual(name = "",
                       values = c("CDFs" = "black",
                                    "Pbox" = "red")) +
    quides (colour =
             guide_legend(override.aes =
                             list(linetype = c(1, 1),
```

```
lwd = c(0.5, 1))) +
        theme(legend.text = element_text(size = 12))
    } else if (values$showpbx) {
      plt <- plt +
        scale_color_manual(name = "",
                           values = c("Pbox" = "red")) +
        theme(legend.text = element_text(size = 12))
    \} else if (values$showcdf)\{
      plt <- plt +
        scale_color_manual(name = "",
                            values = c("CDFs" = "black")) +
        theme(legend.text = element_text(size = 12))
    values$plot <- plt</pre>
    plt
   })
  output$extractTypeInput <- renderUI({</pre>
    if (input$extractInt == TRUE){
      radioButtons(inputId = "extractType",
                   label = "Interval Type: ",
                   choices = c("SRQ Interval",
                                "Probability Interval"))
  })
  output$valInput <- renderUI({</pre>
    if (input$extractInt == TRUE) {
      if (input$extractType == "Probability Interval"){
        lbl = "SRQ Input Value"
        val = round(quantile(values$data[,1], 0.1), 0)
      } else {
        lbl = "Probability Input Value"
        val = 0.95
      numericInput(inputId = "valIn",
                   label = lbl,
value = val)
  })
  output$fileInput <- renderUI({</pre>
    if (input$radioDataSource == "File Upload"){
      fileInput(inputId = "fileIn",
                label = "Upload .csv File")
  })
  # help from https://stackoverflow.com/questions
  # /14810409/save-plots-made-in-a-shiny-app
  output$savePng <- downloadHandler(</pre>
    filename = function(){
      paste(input$pngName,
            ".png",
            sep="")
    },
    content = function(file){
      ggsave(file,
             plot = values$plot,
             device = "png",
             width = 12)
})
```