

# Power Analysis App

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## Introduction/Abstract

The goal of this project is to generate a Shiny web app for conducting simple power analyses on user-provided data. Most online resources currently require users to provide pre-processed data in the form of individual and group averages. Excel spreadsheets with embedded lookup tables for calculating power analyses require users to be comfortable with several advanced excel features, and assumes familiarity with lookup tables such as those provided by Bausell and Li (2002).

These requirements may act as a barrier for some researchers. The overall goal of this project is to encourage researchers to conduct power analyses on pilot data by providing a user-friendly web-based application that takes a csv file with raw data as input, and calculates sample sizes for specified power/alpha levels over a set of statistical tests. As much as possible, the results provided by the script have been cross-validating using G\*Power.

## Dependencies

The shiny script is located in the file `Power_Analysis_App.R`, which in turn calls `ComputeSampleSize.R` for the actual calculations. Taken together, the app uses several libraries to calculate effect sizes and power analyses for the separate statistical tests.

Key amongst these is the WebPower library, which handles the power analyses. For background/additional references relating to the WebPower library, the manual is available [here](#).

```
library(shiny)
library(WebPower)
library(reshape2)
library(effsize)
library(tidyverse)
library(readxl)
library(DescTools)
library(powerAnalysis)
library(shinythemes)
library(rmarkdown)
library(openxlsx)
library(knitr)
library(xtable)
source("Server/ComputeSampleSize.R")
```

## Define UI

The user interface was designed to take either an excel file or a .csv file input containing raw data that will act as pilot data for calculating an effect size. The effect size will then be used to calculate the required sample size, given selected alpha and power levels. Users can select the alpha/power levels they desire; defaults are set to an alpha level of 0.01 and a power level of 0.9.

```

ui <- fluidPage(
  theme = shinytheme("darkly"),

  # App title
  titlePanel("Sample Size Calculator"),

  # Sidebar layout with input and output definitions
  fluidRow(

    ## Sidebar panel for inputs
    #sidebarPanel(
      source("UI/sidebar-ui.R", local = TRUE)$value,

      # Main panel for displaying outputs
      source("UI/mainPanel-ui.R", local = TRUE)$value
    ),

    # Application Instructions
    fluidRow(
      source("UI/appInstructions-ui.R", local = TRUE)$value
    ),

    # About this App (Static)
    fluidRow(
      source("UI/appNotes-ui.R", local = TRUE)$value
    )
  )
)

```

## UI Sidebar

Source = UI/sidebar-ui.R

```

column(4,
  wellPanel(
    # Input: Select a file
    div(h4("Data Input"),
      class = "text-primary",
      fileInput("loadfile",
        "Load an input file (Excel or.csv)",
        multiple = FALSE),
      p("For file formatting tips
        and examples, see Application Instructions
        section, below.",
        class = "text-muted"),
      style = "padding-top: 0px;
        padding-bottom: 0px;"
    ),

  wellPanel(
    ## Input: Select a statistical test
    div(h4("Analysis Options"),
      class = "text-primary"
    ),

```

```

selectInput("test_type",
  "Type of Statistical Test:",
  choices = c("Unpaired T-test",
    "Paired T-test",
    "Chi-squared",
    "One-way ANOVA",
    "Two-way ANOVA"),
  selected = "Unpaired T-test"),

## Input: Select an alpha level
radioButtons("alpha",
  "Alpha Level",
  choices = c(0.05, 0.01),
  selected = 0.01),

## Input: Select a power level
radioButtons("power",
  "Power Level",
  choices = c(0.80, 0.90, 0.95),
  selected = 0.90),
style = "padding-top: 0px;
padding-bottom: 0px;"
))

```

## UI Main Panel

Source = UI/mainPanel-ui.R

```

# Main panel for displaying outputs
column(8,
  conditionalPanel(condition = "output.sample_size",
    wellPanel(
      ## Output: Test type
      tags$h3(textOutput("test_type"),
        class = "text-warning"),

      ## Outout: Results table with
      # sample size calculation
      tableOutput("sample_size"),

      ## Output: Notes re: power analysis
      tags$h4("Analysis Notes",
        class = "text-primary"),
      htmlOutput("test_notes"),

      ## Output: Explaining the variables
      tags$h4("Reported Variables",
        class = "text-primary"),
      uiOutput("variables"),

      ## Alternative Download
      downloadButton(
        "generate_excel_report",
        "Create Excel Report",

```

```

        class = "btn btn-primary"
      )
    )

),
## Error Display
conditionalPanel(condition = "output.error",
  wellPanel(
    # Output: display error codes
    htmlOutput("error")))
)

```

## UI Application Instructions

Source = UI/appInstructions-ui.R

```

column(12,
  wellPanel(
    h3("Application Instructions",
      class = "text-primary"),
    p("To begin, load an Excel or a .csv file containing pilot data.
      Then select the desired statistical
      test using the pull down menu.
      You may also select the desired
      alpha and power levels.
      The suggested default is an alpha
      of 0.01 and a power of 0.9."),
    h4("Formatting your input files",
      class = "text-primary"),
    p("Warning, the .csv format is often finicky. Make sure
      that your file does not have extraneous
      'empty' data columns."),
    h5("T-tests",
      class = "text-warning"),
    p("Data should be separated into two
      adjacent columns (e.g. control vs experimental)."),
    tags$ul(
      tags$li("For unpaired tests,
        the n per condition does
        not have to be equal."),
      tags$li("For paired tests, the number of rows
        must either match;
        uneven rows will be ignored.")
    ),
    h5("Chi-squared tests",
      class="text-warning"),
    p("Data (proportions) should be
      separated into two adjacent columns."),
    h5("One-way ANOVA",
      class = "text-warning"),
    p("Data should be separated into at least 3
      adjacent columns. The number of

```

```

        rows does not have to be even
        across all conditions."),
h5("Two-way ANOVA",
    class = "text-warning"),
p("The power analysis for both main
    effects and the interaction
    effect will be calculated.
    Data should be arranged in columns as follows:"),
tags$ul(
  tags$li("Column 1: Factor A Condition 1"),
  tags$li("Column 2: Factor A Condition 2"),
  tags$li("Column 3: Factor B Condition 1"),
  tags$li("Column 4: Factor B Condition 2")
),
fluidRow(
  column(6,
    div(selectInput("example_type",
      "Download example .csv file:",
      choices = c("Unpaired T-test",
        "Paired T-test",
        "Chi-squared",
        "One-way ANOVA",
        "Two-way ANOVA"),
      selected = "Unpaired T-test"),
      class = "text-warning",
      style = "padding-bottom: 0px"),
    div(downloadLink('downloadData',
      'Download'),
      style = "text-align: right;
        padding-top: 0px"))),
  style = "padding-top: 0px;
    padding-bottom: 5px;")
)

```

## UI Static Application Notes

Source = UI/appNotes-ui.R

```

column(12,
  wellPanel(
    h4("About this application",
      class = "text-primary"),
    p("This Shiny app calculates statistical power analyses
      on user-provided data pilot data. The app takes a .csv
      file with raw data as input, and calculates sample
      sizes for specified power and alpha levels for several
      common statistical tests.", class = "text-muted"),
    p("For all these tests, the assumption is made that the
      data are pulled from a normal distribution,
      i.e. that the statistical test used will be parametric.
      Keep in mind that sample sizes provided may be an
      underestimation in the case where the intention is
      to use non-parametric statistical tests.
      The Prism User Guide suggests that in the absence of

```

```

        easy-to-apply mathematical tools for conducting power
        analyses of non-parametric data, ",
a(href = "https://www.graphpad.com/guides/prism/7/
    statistics/stat_sample_size_for_nonparametric_.htm",
    "values can be estimated calculating the sample size
    for a parametric test and adding 15%",
    .noWS = "outside"),".",
.noWS = c("after-begin", "before-end"),
class = "text-muted"),
p("These calculations are primarily powered by the
    WebPower Library. For background and/or additional
    references relating to the WebPower library, ",
a(href = "https://webpower.psychstat.org/wiki/",
    "a manual and a wiki site are available online",
    .noWS = "outside"),'.',
.noWS = c("after-begin", "before-end"),
class = "text-muted"
),
p("Created by Astra S, Bryant, PhD",
    class = "text-muted"),
style = "padding-top: 0px;
        padding-bottom: 0px;")
)

```

## Define Server Logic / Run the App

The server logic section calls a separate R script containing the sample size calculator (see below), and provides text/table output to the UI. The final line runs the app.

```

server <- function (input, output){
  # Vals will contain all output variables
  vals <- reactiveValues(o = NULL, v = NULL, t = NULL,
    n = NULL, dat = NULL, inputs = NULL)

  # Sample Size Calculations
  dataOutput <- reactive({
    req(input$loadfile) ## Don't run the code unless a file has been selected
    filename <-(input$loadfile$datapath)
    split_filename<-SplitPath(input$loadfile$name)

    ## Import Data, either an xls/xlsx or a csv file
    validate(need(split_filename$extension == "xls" ||
      split_filename$extension == "xlsx" ||
      split_filename$extension == "csv",
      message = "Please select an Excel or .csv file"))

    if (split_filename$extension == "xls" ||
      split_filename$extension == "xlsx"){
      dat <- read_excel(filename)
    } else if (split_filename$extension == "csv") {
      dat <- read.csv(filename)}

    vals$inputs <- split_filename$fullfilename
    vals$dat <- dat
  })
}

```

```

        result<-ComputeSampleSize(dat, input, vals)
    })

    # Parse outputs to Shiny UI
    source("Server/shinyOutputs-srv.R", local = TRUE)

    # Explanations of Reported Variables
    source("Server/variableCb-srv.R", local = TRUE)

    # Generate and Download report
    source("Server/excel-srv.R", local = TRUE)

    # Error Messages
    source("Server/error-srv.R", local = TRUE)

    # Generate example .csv files for download
    source("Server/exampleCSV-srv.R", local = TRUE)
}
shinyApp(ui = ui, server = server)

```

## Parse Outputs to Shiny UI

Source = Server/shinyOutputs-srv.R

```

# Parsing results of sample size
# calculation for Shiny display
output$sample_size<- renderTable({
  result<-dataOutput()
  if (!is.null(result)){
    vals$o <- dplyr::select(result,
                           -c(note,method,url))

    vals$o}
})

outputOptions(output, 'sample_size', suspendWhenHidden = FALSE)

# Parsing statistical test type for Shiny display
output$test_type <- renderText({
  result<-dataOutput()
  if (!is.null(result)){
    vals$t <- dplyr::pull(result,method)%>%
      dplyr::first()
    vals$t
  }
})

# Parsing notes relevant to statistical test type
# for Shiny display
output$test_notes <- renderUI({
  result<-dataOutput()
  if (!is.null(result)){
    #str0 <- c('<h4 class = "text-primary">Notes</h4>')
    str0 <- dplyr::pull(result,note) %>%
      dplyr::first() %>%

```

```

    paste(".", sep = "")

    str1 <- paste('This calculation assumes that data are ',
                  'pulled from a normal distribution.',
                  sep = "")
    str2 <- paste('If you plan to use a non-parametric test, ',
                  'add 15% to the calculated n.', sep = "")
    vals$n <- data.frame(notes = c(str0, str1, str2))
    apply(vals$n, 1, function(x) tags$p(x['notes']))

  }
})

```

## Generate a “Codebook” of Returned Variables

Source = Server/variableCb-srv.R

```

# Variable Codebook
# Explanations of Reported Variables
output$variables <- renderUI({
  result<-dataOutput()
  if (!is.null(result)){
    method <- dplyr::pull(result, method)
    str0 <- c('n = sample size')
    str2 <- c('alpha = significance level (aka false positive rate)')
    str3 <- c('power = statistical power (aka 1 - false negative rate)')

    if (grepl('t-test', method[1])){
      str1<- c('d = effect size (Cohens D)')
      str4<- c('alternative = direction of the alternative hypothesis')
      str5<- c(' ')
      str6<- c(' ')

    } else if (grepl('proportion', method[1])){
      str1<- c('h = effect size')
      str4<- c(' ')
      str5<- c(' ')
      str6<- c(' ')

    } else if (grepl('One-way', method[1])){
      str1<- c('f = effect size (f-ratio)')
      str4<- c('k = number of groups')
      str5<- c(' ')
      str6<- c(' ')

    } else if (grepl('Two-way', method[1])){
      str1<- c('f = effect size (f-ratio)')
      str4<- c('ndf = numerator degrees of freedom')
      str5<- c('ddf = denominator degrees of freedom')
      str6<- c('ng = number of groups')
    }
    vals$v <- data.frame(variables = c(str0, str1, str2, str3, str4, str5, str6))
    apply(vals$v, 1, function(x) tags$p(x['variables']))
  }
})

```



```
}})
```

## Generate Error Messages

Source = Server/error-srv.R

```
# Error Messages
output$error <- renderUI({
  result<-dataOutput()
  if (is.null(result)){
    str0 <-c('<h3 class = "text-danger">Warning</h3>')
    str1 <-c('<p>Number of data columns or rows does
              not match the selected statistical test. </p>
              <p>Please pick another file.</p>')
    str2 <-c('<p class = "text-muted">
              Instructions for correct formatting of .csv files
              can be found under the Application Instructions
              section below. </p>')

    HTML(paste(str0,str1, str2,sep = '<br/>'))
  }
})
outputOptions(output, 'error', suspendWhenHidden = FALSE)
```

## Generate Example .csv Files

These can be downloaded from a handle in the static AppNotes section Source = Server/exampleCSV-srv.R

```
# Generate example .csv files for download
exampleOutput <- reactive({
  req(input$example_type) ## Don't run the code unless an output type
  if (grepl('T-test',input$example_type)){
    example_data <- cbind(Control = c(0.3, 0.2, 0.5),
                          Experimental = c(0.8, 0.7, 1.1))
  } else if (grepl('Chi',input$example_type)){
    example_data <- cbind(Control = 0.5, Experimental = 0.8)
  }else if (grepl('One-way',input$example_type)){
    example_data <- cbind(Control = c(0.3, 0.2, 0.5),
                          ExperimentalA = c(0.8, 0.7, 1.1),
                          ExperimentalB = c(0.5, 0.9, 1.3))
  }else if (grepl('Two-way',input$example_type)){
    example_data <- cbind(Control_ConditionA = c(0.3, 0.2, 0.5),
                          Control_ConditionB = c(0.8, 0.7, 1.1),
                          Experimental_ConditionA = c(0.1, 0.2, 1.1),
                          Experimental_ConditionB = c(0.5, 0.9, 1.3))
  }
})

output$downloadData<- downloadHandler(
  filename = function(){
    paste('example_',input$example_type, '.csv', sep='')
  },
  content = function(con) {
    example_data <- exampleOutput()
    write.csv(example_data,con)
  }
})
```

```
}  
)
```

## Sample Size Calculator

The sample size is calculated in a separate script `ComputeSampleSize.R`.

### Inputs/Outputs

**Input** from the UI includes the raw data, and the user-selected statistical test type, alpha level, and power level.

**Output** to the UI includes the sample size for the desired statistical test, as well as the effect size, alpha level, power level, and notes about whether the given sample sizes refer to the number within a group.

Users may download an excel report containing the outputs and a copy of the inputs.

### Types of Power Calculations

The app will calculate the required sample sizes for 5 different types of statistical tests:

- Unpaired T-test
- Paired T-test
- Chi-squared Test
- One-way ANOVA
- Two-way ANOVA

For all these tests, the assumption is made that the data are pulled from a normal distribution, i.e. that the statistical test used will be parametric. User's should keep in mind that sample sizes provided may be an underestimation in the case where the intention is to use non-parametric statistical tests. The Prism User Guide suggests that in the absence of easy-to-apply mathematical tools for conducting power analyses of non-parametric data, values can be estimated calculating the sample size for a parametric test and adding 15%.

### Unpaired T-test

```
if (input$test_type == "Unpaired T-test"){  
  if (ncol(dat) > 2){return(NULL)}  
  if (nrow(dat) < 2){return(NULL)}  
  dat_melt <- melt(dat,  
    variable.name = 'condition',  
    value.name = 'result',  
    na.rm = TRUE)  
  
  ES <- cohen.d(d = dat_melt$result, f = dat_melt$condition)  
  
  Sample_size <- wp.t(d = ES$estimate,  
    alpha = as.numeric(input$alpha),  
    power = as.numeric(input$power),  
    type = "two.sample")%>%  
    unclass() %>%  
    as_tibble  
  #Sample_size$note <- paste("Note:", Sample_size$note)  
  return(Sample_size)
```

## Paired T-test

```
} else if (input$test_type == "Paired T-test"){
  if (ncol(dat) > 2){return(NULL)}
  if (nrow(dat) < 2){return(NULL)}
  dat_complete <- dat[complete.cases(dat),]
  n <- nrow(dat_complete)
  dat_complete$id <- (1:n)

  dat_melt <- reshape2::melt(dat_complete,
    measure.vars = 1:2,
    variable.name = 'condition',
    value.name = 'result',
    na.rm = TRUE)

  ES <- cohen.d(d = dat_melt$result,
    f = dat_melt$condition,
    subject = dat_melt$id,
    paired = TRUE)

  Sample_size <- wp.t(d = ES$estimate,
    alpha = as.numeric(input$alpha),
    power = as.numeric(input$power),
    type = "paired") %>%
    unclass() %>%
    as_tibble
  #Sample_size$note <- paste("Note:", Sample_size$note)
  return(Sample_size)
```

## Chi-squared Test

```
} else if (input$test_type == "Chi-squared"){
  if (ncol(dat) > 2){return(NULL)}
  if (nrow(dat) > 2){return(NULL)}
  dat_melt <- melt(dat,
    variable.name = 'condition',
    value.name = 'proportion')
  ES <- (2 * asin(sqrt(dat_melt$proportion[[1]]))) -
    (2 * asin(sqrt(dat_melt$proportion[[2]])))
  Sample_size <- wp.prop(h = ES,
    alpha = 0.01, ##as.numeric(input$alpha),
    power = 0.9, ##as.numeric(input$power),
    type = "2p") %>%
    unclass() %>%
    as_tibble
  # Sample_size$note <- paste("Note:", Sample_size$note)
  Sample_size <- Sample_size %>%
    select("n", "h", everything())
  return(Sample_size)
```

## One-way ANOVA

```

} else if (input$test_type == "One-way ANOVA"){
  if (ncol(dat) < 3){return(NULL)}
  k = ncol(dat) # Number of groups

  dat_melt <- melt(dat,
    variable.name = 'condition',
    value.name = 'result',
    na.rm = TRUE
  )

  anova1 <- aov(formula = dat_melt$result ~ dat_melt$condition)
  summary(anova1)
  etasquared <- EtaSq(anova1)
  CohensF <- sqrt(etasquared[2]/(1-etasquared[2]))
  Sample_size <- wp.anova(k = k,
    f = CohensF,
    alpha = as.numeric(input$alpha),
    power = as.numeric(input$power)) %>%
    unclass() %>%
    as_tibble

  Sample_size$n <- Sample_size$n/Sample_size$k
  Sample_size$note <- c("n is the sample size *in each group*")
  Sample_size$method <- c("One-way ANOVA")
  Sample_size <- Sample_size %>%
    dplyr::select("n", "f", everything())
  return(Sample_size)
}

```

## Two-way ANOVA

```

} else if (input$test_type == "Two-way ANOVA"){
  if (ncol(dat) < 4){return(NULL)}
  k <- ncol(dat) # Total number of groups
  r_num <- 2 ## Number of factors located in rows,
  ## for a 2 way anova this is hard-wired
  c_num <- k/r_num ## Number of factors located in columns

  ## Next couple of lines thanks of K. Zalocusky, PhD
  id_list <- c(1:r_num)
  id_list2 <- c(1:c_num)

  ids <- expand.grid(id_list, id_list2)
  ids <- ids[rep(seq_len(nrow(ids)), each = nrow(dat)), ]

  dat_melt <- melt(dat)
  dat_melt <- cbind(dat_melt, ids)
  dat_melt <- dat_melt[!is.na(dat_melt$value),] ## Get rid of NA values

  dat_melt$Var1 <- as_factor(dat_melt$Var1)
  dat_melt$Var2 <- as_factor(dat_melt$Var2)

  anova2 <- aov(formula = dat_melt$value ~ dat_melt$Var1 * dat_melt$Var2)
  etasquared <- EtaSq(anova2)
}

```

```

CohensF <- sqrt(etasquared[,2]/(1-etasquared[,2]))
ndf <- summary.aov(anova2) %>%
  flatten_df() %>%
  dplyr::select(Df) %>%
  pull()

Sample_size_Interaction <- wp.kanova(f = CohensF[
  "dat_melt$Var1:dat_melt$Var2"],
  ndf = ndf[3],
  alpha = as.numeric(input$alpha),
  power = as.numeric(input$power),
  ng = 4) %>%
  unclass() %>%
  as_tibble %>%
  add_column(name = 'Interaction Effect', .before = "n")

Sample_size_Var1 <- wp.kanova(f = CohensF["dat_melt$Var1"],
  ndf = ndf[1],
  alpha = as.numeric(input$alpha),
  power = as.numeric(input$power),
  ng = 4) %>%
  unclass() %>%
  as_tibble %>%
  add_column(name = 'Main Effect of Rows', .before = "n")

Sample_size_Var2 <- wp.kanova(f = CohensF["dat_melt$Var2"],
  ndf = ndf[2],
  alpha = as.numeric(input$alpha),
  power = as.numeric(input$power),
  ng = 4) %>%
  unclass() %>%
  as_tibble() %>%
  add_column(name = 'Main Effect of Cols', .before = "n")

Sample_size <- full_join(x = Sample_size_Var1,y = Sample_size_Var2) %>%
  full_join(y = Sample_size_Interaction)

Sample_size$n <- Sample_size$n/Sample_size$ng
Sample_size$note <- c("n is the sample size *in each group*")
Sample_size$method <- c("Two-way ANOVA")
Sample_size <- Sample_size %>%
  dplyr::select("name", "n", "f", everything())
return(Sample_size)
}

```