

Histopathology Research Template

true

2019-11-08

Contents

1	Define Variable Types	3
2	Overview / Exploratory Data Analysis (EDA)	3
3	Information value	3
4	Feature imbalance	3
5	Memory usage	3
6	SmartEDA	3
6.1	Default ExpParcoord funciton	3
6.2	With Stratified rows and selected columns only	3
6.3	Numeric variable summary	5
6.4	Adding filters for complete data (like base Subset)	5
6.5	Filter unique value from all the numeric variables	5
6.6	Adding filters at variable level	5
7	impute	6
7.1	impute continious	6
7.2	impute categorical	6
7.3	impute outlier	6
8	transform	6
8.1	min -max	6
8.2	skewness	6
8.3	log	6
9	binning	6
9.1	optimal binning	6
9.2	standardize	6
10	data transformation report	6
11	inspectdf	6
12	Descriptive Statistics	7
12.1	Table 1	7
12.2	Categorical Variables	7
12.2.1	Split-Group Stats Categorical	7
12.2.2	Grouped Categorical	7
12.3	Continious Variables	8
12.3.1	Split-Group Stats Continious	8
12.3.2	Grouped Continious	8
13	Cross Tables	9

14 Plots	10
14.1 Categorical Variables	10
15 Plots	10
15.1 Continious Variables	10
16 Hypothesis Tests	10
16.1 Tests of Normality	10
16.2 Categorical	10
16.2.1 Chi-Square Cramer Association Predictive Power	10
16.3 Continious	10
16.4 Odds	10
17 ROC	11
18 Decision Tree	11
19 Survival Analysis	11
20 Pairwise comparison	11
21 Multivariate Analysis Survival	11
22 KM plot	11
23 Separate tables	12
24 Pipe together	12
25 Separate tables	13
26 Pipe together	13
27 OR plot	13
28 HR plot (not fully tested)	13
29 ANOVA	13
30 Save Final Data	17
31 Final Data Summary	18
32 Software and Libraries Used	19
32.1 data[order(data\$References),]	19
33 Session Info	22
34 Notes	25
<pre>knitr::opts_chunk\$set(echo = TRUE, #change to TRUE eval = TRUE, message = FALSE, warning = FALSE, comment = NA, tidy = TRUE,</pre>	

```

    fig.path = here::here("figs/")
  )

# https://cran.r-project.org/web/packages/exploreR/vignettes/exploreR.html
# exploreR::reset()

```

1 Define Variable Types

2 Overview / Exploratory Data Analysis (EDA)

3 Information value

4 Feature imbalance

5 Memory usage

6 SmartEDA

Create HTML EDA report

Create a exploratory data analysis report in HTML format

```
ExpReport(Carseats,Target="Urban",label=NULL,theme="Default",op_file="test.html",op_dir=getwd(),sc=2,
sn=2,Rc="Yes")
```

Quantile-quantile plot for numeric variables

```
ExpOutQQ(CData,nlim=10,fname=NULL,Page=c(2,2),sample=4)
```

Parallel Co-ordinate plots

6.1 Default ExpParcoord function

```
ExpParcoord(CData,Group=NULL,Stsize=NULL,Nvar=c("Price","Income","Advertising","Population","Age",
"Education"))
```

6.2 With Stratified rows and selected columns only

```
ExpParcoord(CData,Group="ShelveLoc",Stsize=c(10,15,20),Nvar=c("Price","Income"),Cvar=c("Ur-
ban","US")) ## Without stratification ExpParcoord(CData,Group="ShelveLoc",Nvar=c("Price","In-
come"),Cvar=c("Urban","US"),scale=NULL)
```

Exploratory analysis - Custom tables, summary statistics

Descriptive summary on all input variables for each level/combination of group variable. Also while running the analysis we can filter row/cases of the data.

```
ExpCustomStat(Carseats,Cvar=c("US","Urban","ShelveLoc"),goby=FALSE)
ExpCustomStat(Carseats,Cvar=c("US","Urban"),goby=TRUE,filt=NULL)
ExpCustomStat(Carseats,Cvar=c("US","Urban","ShelveLoc"),goby=TRUE,filt=NULL)
ExpCustomStat(Carseats,Cvar=c("US","Urban"),goby=TRUE,filt="Population>150")
ExpCustomStat(Carseats,Cvar=c("US","ShelveLoc"),goby=TRUE,filt="Urban=='Yes' & Population>150")
```

```

ExpCustomStat(Carseats,Cvar=c("US","Urban","ShelveLoc","Education"),gby=FALSE)
ExpCTable(Carseats,Target=NULL,clim=5,nlim=15,round=2,bin=NULL,per=F)
ExpCustomStat(Carseats,Cvar=c("US","Urban","ShelveLoc"),gby=FALSE)
ExpCustomStat(Carseats,Cvar=c("US","Urban"),gby=TRUE,flt=NULL)
ExpCustomStat(Carseats,Cvar=c("US","Urban","ShelveLoc"),gby=TRUE,flt=NULL)
ExpCustomStat(Carseats,Cvar=c("US","Urban"),gby=TRUE,flt="Population>150")
ExpCustomStat(Carseats,Cvar=c("US","ShelveLoc"),gby=TRUE,flt="Urban=='Yes' & Population>150")

options(width = 150) ExpCustomStat(Carseats,Nvar=c("Population","Sales","CompPrice","Income"),stat
= c('Count','mean','sum','var','sd','min','max','IQR'))

ExpCustomStat(Carseats,Nvar=c("Population","Sales","CompPrice","Income"),stat = c('min','p0.25','median','p0.75','max'))

options(width = 150) ExpCustomStat(Carseats,Nvar=c("Population","Sales","CompPrice","Income"),stat
= c('Count','mean','sum','var','min','median','max'),flt="Urban=='Yes' ")

options(width=150) ExpCustomStat(Carseats,Nvar=c("Population","Sales","CompPrice","Income"),stat =
c('Count','mean','sum','median','IQR'),flt="Urban=='Yes' & Population>150")

data_sam = Carseats[,] data_sam[sample(1:400,30),"Sales"] <- 999 data_sam[sample(1:400,20),"Comp
Price"] <- -9 data_sam[sample(1:400,45),"Income"] <- 999 ExpCustomStat(data_sam,Nvar=c("Popula
tion","Sales","CompPrice","Income"),stat = c('Count','mean','sum','min'),flt="All %ni% c(999,-9)")

options(width = 150) ExpCustomStat(Carseats,Nvar=c("Population","Sales","CompPrice","Educa
tion","Income"),stat = c('Count','mean','sum','var','sd','IQR','median'),flt=c("ShelveLoc=='Good'Urban=='Yes'Price>=150"))

options(width = 150) ExpCustomStat(Carseats,Cvar = c("Urban","ShelveLoc"), Nvar=c("Popula
tion","Sales"), stat = c('Count','Prop','mean','min','P0.25','median','p0.75','max'),gby=FALSE)

options(width = 150) ExpCustomStat(Carseats,Cvar = c("Urban","US","ShelveLoc"), Nvar=c("Comp
Price","Income"), stat = c('Count','Prop','mean','sum','PS','min','max','IQR','sd'), gby = TRUE)

options(width = 150) ExpCustomStat(Carseats,Cvar = c("Urban","US","ShelveLoc"), Nvar=c("Comp
Price","Income"), stat = c('Count','Prop','mean','sum','PS','median','IQR'), gby = TRUE,flt="Ur
ban=='Yes' ")

options(width = 150) data_sam = Carseats[,] data_sam[sample(1:400,30),"Sales"] <- 888 data_sam[sam
ple(1:400,20),"CompPrice"] <- 999 data_sam[sample(1:400,45),"Income"] <- 999 ExpCustomStat(data_sam,Cvar
= c("Urban","US","ShelveLoc"), Nvar=c("Sales","CompPrice","Income"), stat = c('Count','Prop','mean','sum','PS'),
gby = TRUE,flt="All %ni% c(888,999)")

ExpCustomStat(Carseats,Cvar = c("Urban","US"), Nvar=c("Population","Sales","CompPrice"), stat =
c('Count','Prop','mean','sum','var','IQR'), flt=c("ShelveLoc=='Good'Urban=='Yes'Price>=150"))

options(width = 150) ExpCustomStat(Carseats,Cvar = c("Urban"), Nvar=c("Population","Sales"), stat =
c('Count','Prop'),gby=TRUE,dcast=TRUE)

##Frequency table for categorical variables ExpCustomStat(Carseats,Cvar=c("US","Urban","Shelve
Loc"),gby=FALSE)

##Crosstabulation between categorical variables ExpCustomStat(Carseats,Cvar=c("US","Urban"),gby=TRUE,flt=NULL)
ExpCustomStat(Carseats,Cvar=c("US","Urban","ShelveLoc"),gby=TRUE,flt=NULL)

##Adding filters for custom tables ExpCustomStat(Carseats,Cvar=c("US","Urban"),gby=TRUE,flt="Pop
ulation>150") ExpCustomStat(Carseats,Cvar=c("US","ShelveLoc"),gby=TRUE,flt="Urban=='Yes' &
Population>150")

```

6.3 Numeric variable summary

```
ExpCustomStat(Carseats,Nvar=c("Population","Sales","CompPrice","Income"),stat = c('Count','mean','sum','var','min','median','p0.25','p0.75','max'))
ExpCustomStat(Carseats,Nvar=c("Population","Sales","CompPrice","Income"),stat = c('min','p0.25','median','p0.75','max'))
```

6.4 Adding filters for complete data (like base Subset)

```
ExpCustomStat(Carseats,Nvar=c("Population","Sales","CompPrice","Income"),stat = c('Count','mean','sum','var'),filt="Urban=='Yes'")
ExpCustomStat(Carseats,Nvar=c("Population","Sales","CompPrice","Income"),stat = c('Count','mean','sum'),filt="Urban=='Yes' & Population>150")
```

6.5 Filter unique value from all the numeric variables

```
ExpCustomStat(data_sam,Nvar=c("Population","Sales","CompPrice","Income"),stat = c('Count','mean','sum','min'),filt="%ni% c(999,-9)")
```

6.6 Adding filters at variable level

```
ExpCustomStat(Carseats,Nvar=c("Population","Sales","CompPrice","Education","Income"),stat = c('Count','mean','sum','var','sd','IQR','median'),filt=c("ShelveLoc=='Good'Urban=='Yes'Price>=150^~US=='Yes'"))
```

```
##Numerical summaries by category ##Variable summary report (One group variable) ExpCustomStat(Carseats,Cvar = c("Urban","ShelveLoc"), Nvar=c("Population","Sales"), stat = c('Count','Prop','mean','min','P0.25','median','p0.75','max'),gby=FALSE)
```

```
##Variable summary report (More than One group variable) ExpCustomStat(Carseats,Cvar = c("Urban","US","ShelveLoc"), Nvar=c("CompPrice","Income"), stat = c('Count','Prop','mean','sum','PS','min','max','IQR','sd'),gby = TRUE)
```

```
##Variable summary report (More than One group variable) with filter ExpCustomStat(Carseats,Cvar = c("Urban","US","ShelveLoc"), Nvar=c("CompPrice","Income"), stat = c('Count','Prop','mean','sum','PS','P0.25','median','p0.75'), gby = TRUE,filt="Urban=='Yes'") ExpCustomStat(data_sam,Cvar = c("Urban","US","ShelveLoc"), Nvar=c("Sales","CompPrice","Income"), stat = c('Count','Prop','mean','sum','PS'), gby = TRUE,filt="All %ni% c(888,999)") ExpCustomStat(Carseats,Cvar = c("Urban","US"), Nvar=c("Population","Sales","CompPrice"), stat = c('Count','Prop','mean','sum','var','min','max'), filt=c("ShelveLoc=='Good'Urban=='Yes'"))
```

```
iris %>% mutate(sumVar = rowSums(.[1:4]))
```

```
iris %>% mutate(sumVar = rowSums(select(., contains("Sepal")))) %>% head
```

```
iris %>% mutate(sumVar = select(., contains("Sepal"))) %>% rowSums() %>% head
```

```
iRenameColumn.R
```

```
iSelectColumn.R
```

```
<= 22 Low
```

```
>= 23 & <= 41 Average
```

```
>=42 High
```

7 impute

7.1 impute continuous

7.2 impute categorical

7.3 impute outlier

8 transform

8.1 min -max

8.2 skewness

8.3 log

9 binning

9.1 optimal binning

9.2 standardize

10 data transformation report

11 inspectdf

12 Descriptive Statistics

12.1 Table 1

Report Data

12.2 Categorical Variables

12.2.1 Split-Group Stats Categorical

12.2.2 Grouped Categorical

12.3 Continious Variables

12.3.1 Split-Group Stats Continious

12.3.2 Grouped Continious

13 Cross Tables

14 Plots

14.1 Categorical Variables

15 Plots

15.1 Continious Variables

16 Hypothesis Tests

16.1 Tests of Normality

16.2 Categorical

16.2.1 Chi-Square Cramer Association Predictive Power

16.3 Continious

16.4 Odds

17 ROC

18 Decision Tree

19 Survival Analysis

20 Pairwise comparison

21 Multivariate Analysis Survival



22 KM plot

```
explanatory = c("perfor.factor") dependent = "Surv(time, status)" colon_s %>% surv.plot(dependent, explanatory, xlab="Time (days)",  
pval=TRUE, legend="none")
```

Notes

Use `Hmisc::label()` to assign labels to variables for tables and plots.

```
label(colon_s$age.factor) = "Age (years)"
```

Export dataframe tables directly or to R Markdown using `knitr::kable()`.

Note wrapper `summary.missing()` can be useful. Wraps `mice::md.pattern`.

```
colon_s %>% summary.missing(dependent, explanatory)
```

Where a multivariable model contains a subset of the variables specified in the full univariable set, this can be specified.

```
explanatory = c("age.factor", "sex.factor", "obstruct.factor", "perfor.factor") explanatory.multi = c("age.factor", "obstruct.factor") dependent = 'mort_5yr' colon_s %>% summarizer(dependent, explanatory, explanatory.multi)
```

Random effects.

```
e.g. lme4::glmer(dependent ~ explanatory + (1 | random_effect), family="binomial")
```

```
explanatory = c("age.factor", "sex.factor", "obstruct.factor", "perfor.factor") explanatory.multi = c("age.factor", "obstruct.factor") random.effect = "hospital" dependent = 'mort_5yr' colon_s %>% summarizer(dependent, explanatory, explanatory.multi, random.effect)
```

metrics=TRUE provides common model metrics.

```
colon_s %>% summarizer(dependent, explanatory, explanatory.multi, metrics=TRUE)
```

Cox proportional hazards

```
e.g. survival::coxph(dependent ~ explanatory)
```

```
explanatory = c("age.factor", "sex.factor", "obstruct.factor", "perfor.factor") dependent = "Surv(time, status)"
```

```
colon_s %>% summarizer(dependent, explanatory)
```

Rather than going all-in-one, any number of subset models can be manually added on to a summary.factorlist() table using summarizer.merge(). This is particularly useful when models take a long-time to run or are complicated.

Note requirement for glm.id=TRUE. fit2df is a subfunction extracting most common models to a dataframe.

```
explanatory = c("age.factor", "sex.factor", "obstruct.factor", "perfor.factor") explanatory.multi = c("age.factor", "obstruct.factor") random.effect = "hospital" dependent = 'mort_5yr'
```

23 Separate tables

```
colon_s %>% summary.factorlist(dependent, explanatory, glm.id=TRUE) -> example.summary
```

```
colon_s %>% glmuni(dependent, explanatory) %>% fit2df(estimate.suffix=" (univariable)") -> example.univariable
```

```
colon_s %>% glmmulti(dependent, explanatory) %>% fit2df(estimate.suffix=" (multivariable)") -> example.multivariable
```

```
colon_s %>% glmmixed(dependent, explanatory, random.effect) %>% fit2df(estimate.suffix=" (multilevel)") -> example.multilevel
```

24 Pipe together

```
example.summary %>% summarizer.merge(example.univariable) %>% summarizer.merge(example.multivariable) %>% summarizer.merge(example.multilevel) %>% select(-c(glm.id, index)) -> example.final
```

Cox Proportional Hazards example with separate tables merged together.

```
explanatory = c("age.factor", "sex.factor", "obstruct.factor", "perfor.factor") explanatory.multi = c("age.factor", "obstruct.factor") dependent = "Surv(time, status)"
```

25 Separate tables

```
colon_s %>% summary.factorlist(dependent, explanatory, glm.id=TRUE) -> example2.summary  
colon_s %>% coxphuni(dependent, explanatory) %>% fit2df(estimate.suffix=" (univariable)") -> exam-  
ple2.univariable  
colon_s %>% coxphmulti(dependent, explanatory.multi) %>% fit2df(estimate.suffix=" (multivariable)") ->  
example2.multivariable
```

26 Pipe together

```
example2.summary %>% summarizer.merge(example2.univariable) %>% summarizer.merge(example2.mul-  
tivariable) %>% select(-c(glm.id, index)) -> example2.final example2.final
```

27 OR plot

```
explanatory = c("age.factor", "sex.factor", "obstruct.factor", "perfor.factor") dependent = 'mort_5yr'  
colon_s %>% or.plot(dependent, explanatory) # Previously fitted models (glmmulti() or glmmixed())  
can be provided directly to glmfit
```

28 HR plot (not fully tested)

```
explanatory = c("age.factor", "sex.factor", "obstruct.factor", "perfor.factor") dependent = "Surv(time, sta-  
tus)" colon_s %>% hr.plot(dependent, explanatory, dependent_label = "Survival") # Previously fitted  
models (coxphmulti) can be provided directly using coxfit
```

29 ANOVA

Some Text ile sağkalım açısından bir ilişki bulunmamıştır ($p = 0.22$).

Some Text ile sağkalım açısından bir ilişki bulunmamıştır ($p = 0.22$).

Some Text

İstatistik Metod:, , Sürekli verilerin ortalama, standart sapma, median, minimum ve, maksimum değerleri verildi. Kategorik veriler ve gruplanan sürekli, veriler için frekans tabloları oluşturuldu. Genel sağkalım, analizinde ölüm tarihi ve son başvuru tarihi hasta dosyalarından, elde edildi. Sağkalım analizinde Kaplan-Meier grafikleri,, Log-rank testi ve Cox-Regresyon testleri uygulandı. Analizler, R-project (version 3.6.0) ve RStudio ile survival ve finalfit, paketleri kullanılarak yapıldı. p değeri 0.05 düzeyinde anlamlı, olarak kabul edildi., , R Core Team (2019). R: A language and environment for statistical, computing. R Foundation for Statistical Computing, Vienna,, Austria. URL <https://www.R-project.org/>., , Therneau T (2015). A Package for Survival Analysis in S. version, 2.38, <https://CRAN.R-project.org/package=survival>, , Terry M. Therneau, Patricia M. Grambsch (2000). Modeling Survival, Data: Extending the Cox Model. Springer, New York. ISBN, 0-387-98784-3., , Ewen Harrison, Tom Drake and Riinu Ots (2019). finalfit: Quickly, Create Elegant Regression Results Tables and Plots when Modelling., R package version 0.9.6. <https://github.com/ewenharrison/finalfit>

İstatistik Metod:, , Sürekli verilerin ortalama, standart sapma, median, minimum ve, maksimum değerleri verildi. Kategorik veriler ve gruplanan sürekli, veriler için frekans tabloları oluşturuldu. Genel sağkalım, analizinde ölüm tarihi ve son başvuru tarihi hasta dosyalarından, elde edildi. Sağkalım analizinde Kaplan-Meier grafikleri,, Log-rank testi ve Cox-Regresyon testleri uygulandı. Analizler, R-project (version 3.6.0) ve RStudio ile survival ve finalfit, paketleri kullanılarak yapıldı. p değeri 0.05 düzeyinde anlamlı, olarak kabul edildi., , R Core Team (2019). R: A language and environment for statistical, computing. R Foundation for Statistical Computing, Vienna,, Austria. URL <https://www.R-project.org/>., , Therneau T (2015). A Package for Survival Analysis in S. version, 2.38, <https://CRAN.R-project.org/package=survival>, , Terry M. Therneau, Patricia M. Grambsch (2000). Modeling Survival, Data: Extending the Cox Model. Springer, New York. ISBN, 0-387-98784-3., , Ewen Harrison, Tom Drake and Riinu Ots (2019). finalfit: Quickly, Create Elegant Regression Results Tables and Plots when Modelling., R package version 0.9.6. <https://github.com/ewenharrison/finalfit>

Text Here

Text Here

30 Save Final Data

saved data after analysis to/Users/serdarbalciold/histopathology-template/data/histopathology-template2019-11-08.xlsx : 2019-11-08 23:38:35

31 Final Data Summary

32 Software and Libraries Used

To cite R in publications use:

R Core Team (2019). R: A language and environment for statistical computing. R Foundation for Statistical Computing, Vienna, Austria. URL <https://www.R-project.org/>.

A BibTeX entry for LaTeX users is

```
@Manual{,  
  title = {R: A Language and Environment for Statistical Computing},  
  author = {{R Core Team}},  
  organization = {R Foundation for Statistical Computing},  
  address = {Vienna, Austria},  
  year = {2019},  
  url = {https://www.R-project.org/},  
}
```

We have invested a lot of time and effort in creating R, please cite it when using it for data analysis. See also 'citation("pkgname")' for citing R packages.

The jamovi project (2019). jamovi. (Version 0.9) [Computer Software]. Retrieved from <https://www.jamovi.org>. R Core Team (2018). R: A Language and environment for statistical computing. [Computer software]. Retrieved from <https://cran.r-project.org/>. Fox, J., & Weisberg, S. (2018). car: Companion to Applied Regression. [R package]. Retrieved from <https://cran.r-project.org/package=car>.

32.1 data[order(data\$References),]

Ewen Harrison, Tom Drake and Riinu Ots (2019). finalfit: Quickly Create Elegant Regression Results Tables and Plots when Modelling. R package version 0.9.6. <https://github.com/ewenharrison/finalfit>
Hadley Wickham and Jennifer Bryan (2019). readxl: Read Excel Files. R package version 1.3.1. <https://CRAN.R-project.org/package=readxl>
Hadley Wickham, Romain François, Lionel Henry and Kirill Müller (2019). dplyr: A Grammar of Data Manipulation. R package version 0.8.3. <https://CRAN.R-project.org/package=dplyr>
Makowski, D. & Lüdtke, D. (2019). The report package for R: Ensuring the use of best practices for results reporting. CRAN. Available from <https://github.com/easystats/report>. doi: .
Patil I (2018). ggstatsplot: 'ggplot2' Based Plots with Statistical Details. doi: 10.5281/zenodo.2074621 (URL:<https://doi.org/10.5281/zenodo.2074621>), <URL:<https://CRAN.R-project.org/package=ggstatsplot>>.
Rinker, T. W. (2018). wakefield: Generate Random Data. version 0.3.3. Buffalo, New York. <https://github.com/trinker/wakefield>
Sam Firke (2019). janitor: Simple Tools for Examining and Cleaning Dirty Data. R package version 1.2.0. <https://CRAN.R-project.org/package=janitor>

To cite package 'tidyverse' in publications use:

Hadley Wickham (2017). tidyverse: Easily Install and Load the 'Tidyverse'. R package version 1.2.1. <https://CRAN.R-project.org/package=tidyverse>

A BibTeX entry for LaTeX users is

```
@Manual{,
  title = {tidyverse: Easily Install and Load the 'Tidyverse'},
  author = {Hadley Wickham},
  year = {2017},
  note = {R package version 1.2.1},
  url = {https://CRAN.R-project.org/package=tidyverse},
}
```

To cite package 'readxl' in publications use:

Hadley Wickham and Jennifer Bryan (2019). readxl: Read Excel Files. R package version 1.3.1.
<https://CRAN.R-project.org/package=readxl>

A BibTeX entry for LaTeX users is

```
@Manual{,
  title = {readxl: Read Excel Files},
  author = {Hadley Wickham and Jennifer Bryan},
  year = {2019},
  note = {R package version 1.3.1},
  url = {https://CRAN.R-project.org/package=readxl},
}
```

To cite package 'janitor' in publications use:

Sam Firke (2019). janitor: Simple Tools for Examining and Cleaning Dirty Data. R package version 1.2.0.
<https://CRAN.R-project.org/package=janitor>

A BibTeX entry for LaTeX users is

```
@Manual{,
  title = {janitor: Simple Tools for Examining and Cleaning Dirty Data},
  author = {Sam Firke},
  year = {2019},
  note = {R package version 1.2.0},
  url = {https://CRAN.R-project.org/package=janitor},
}
```

To cite in publications use:

Makowski, D. & Lüdecke, D. (2019). The report package for R: Ensuring the use of best practices for results reporting. CRAN. Available from <https://github.com/easystats/report>. doi: .

A BibTeX entry for LaTeX users is

```
@Article{,
  title = {The report package for R: Ensuring the use of best practices for results reporting},
  author = {{Makowski} and {Dominique} and {Lüdecke} and {Daniel}},
  journal = {CRAN},
}
```

```

year = {2019},
note = {R package},
url = {https://github.com/easystats/report},
}

```

To cite package 'finalfit' in publications use:

Ewen Harrison, Tom Drake and Riinu Ots (2019). finalfit: Quickly Create Elegant Regression Results Tables and Plots when Modelling. R package version 0.9.6.
<https://github.com/ewenharrison/finalfit>

A BibTeX entry for LaTeX users is

```

@Manual{,
  title = {finalfit: Quickly Create Elegant Regression Results Tables and Plots when
Modelling},
  author = {Ewen Harrison and Tom Drake and Riinu Ots},
  year = {2019},
  note = {R package version 0.9.6},
  url = {https://github.com/ewenharrison/finalfit},
}

```

Patil I (2018). _ggstatsplot: 'ggplot2' Based Plots with Statistical Details_. doi: 10.5281/zenodo.2074621 (URL: <https://doi.org/10.5281/zenodo.2074621>), <URL: <https://CRAN.R-project.org/package=ggstatsplot>>.

A BibTeX entry for LaTeX users is

```

@Manual{,
  title = {ggstatsplot: 'ggplot2' Based Plots with Statistical Details},
  author = {Indrajeet Patil},
  year = {2018},
  url = {https://CRAN.R-project.org/package=ggstatsplot},
  doi = {10.5281/zenodo.2074621},
}

```

33 Session Info

```
R version 3.6.0 (2019-04-26)
Platform: x86_64-apple-darwin15.6.0 (64-bit)
Running under: macOS 10.15.1

Matrix products: default
BLAS:   /Library/Frameworks/R.framework/Versions/3.6/Resources/lib/libRblas.0.dylib
LAPACK: /Library/Frameworks/R.framework/Versions/3.6/Resources/lib/libRlapack.dylib

locale:
[1] en_US.UTF-8/en_US.UTF-8/en_US.UTF-8/C/en_US.UTF-8/en_US.UTF-8

attached base packages:
[1] stats      graphics  grDevices  utils      datasets  methods    base

other attached packages:
[1] wakefield_0.3.3  ggstatsplot_0.1.2 finalfit_0.9.6    report_0.1.0
[5] janitor_1.2.0    readxl_1.3.1      dplyr_0.8.3

loaded via a namespace (and not attached):
 [1] estimability_1.3      SparseM_1.77
 [3] coda_0.19-3           tidyr_1.0.0
 [5] ggplot2_3.2.1         acepack_1.4.1
 [7] knitr_1.25            multcomp_1.4-10
 [9] data.table_1.12.6     rpart_4.1-15
[11] inline_0.3.15         generics_0.0.2
[13] callr_3.3.2           cowplot_1.0.0
[15] TH.data_1.0-10        mice_3.6.0
[17] future_1.14.0         webshot_0.5.1
[19] xml2_1.2.2            httpuv_1.5.2
[21] StanHeaders_2.19.0    assertthat_0.2.1
[23] WRS2_1.0-0            xfun_0.10
[25] hms_0.5.1             evaluate_0.14
[27] promises_1.0.1        DEoptimR_1.0-8
[29] htmlwidgets_1.5.1     mcmc_0.9-6
[31] reshape_0.8.8         stats4_3.6.0
[33] paletteer_0.2.1       purrr_0.3.3
[35] ellipsis_0.3.0        rcompanion_2.3.0
[37] backports_1.1.5       insight_0.6.0
[39] ggcorrplot_0.1.3      MCMCpack_1.4-4
[41] libcoin_1.0-5         jmvcore_1.0.0
[43] vctr_0.2.0            quantreg_5.51
[45] here_0.1              sjlabelled_1.1.1
[47] abind_1.4-5           metaBMA_0.6.2
[49] robustbase_0.93-5     checkmate_1.9.4
[51] emmeans_1.4.1         prettyunits_1.0.2
[53] mnormt_1.5-5          cluster_2.1.0
[55] lazyeval_0.2.2        crayon_1.3.4
[57] pkgconfig_2.0.3       nlme_3.1-141
[59] statsExpressions_0.1.1 nnet_7.3-12
[61] rlang_0.4.1           globals_0.12.4
[63] lifecycle_0.1.0       mitml_0.3-7
[65] miniUI_0.1.1.1        groupedstats_0.0.9
```

[67] skimr_1.0.7	LaplacesDemon_16.1.1
[69] MatrixModels_0.4-1	sandwich_2.5-1
[71] EMT_1.1	modelr_0.1.5
[73] cellranger_1.1.0	rprojroot_1.3-2
[75] matrixStats_0.55.0	broomExtra_0.0.5
[77] lmtest_0.9-37	Matrix_1.2-17
[79] loo_2.1.0	mc2d_0.1-18
[81] carData_3.0-2	boot_1.3-23
[83] zoo_1.8-6	pan_1.6
[85] base64enc_0.1-3	processx_3.4.1
[87] viridisLite_0.3.0	rjson_0.2.20
[89] parameters_0.2.0.1	ggExtra_0.9
[91] stringr_1.4.0	multcompView_0.1-7
[93] coin_1.3-1	robust_0.4-18.1
[95] readr_1.3.1	ggsignif_0.6.0
[97] scales_1.0.0	magrittr_1.5
[99] plyr_1.8.4	compiler_3.6.0
[101] rstantools_2.0.0	kableExtra_1.1.0
[103] RColorBrewer_1.1-2	lme4_1.1-21
[105] rrcov_1.4-7	cli_1.1.0
[107] listenv_0.7.0	pbapply_1.4-2
[109] ps_1.3.0	TMB_1.7.15
[111] Brodingtonag_1.2-6	htmlTable_1.13.2
[113] formatR_1.7	Formula_1.2-3
[115] MASS_7.3-51.4	mgcv_1.8-29
[117] tidyselect_0.2.5	stringi_1.4.3
[119] forcats_0.4.0	yaml_2.2.0
[121] latticeExtra_0.6-28	ggrepel_0.8.1
[123] bridgesampling_0.7-2	grid_3.6.0
[125] manipulate_1.0.1	tools_3.6.0
[127] parallel_3.6.0	rio_0.5.16
[129] rstudioapi_0.10	foreign_0.8-72
[131] gridExtra_2.3	pairwiseComparisons_0.1.1
[133] digest_0.6.22	shiny_1.3.2
[135] explore_0.5.1	nortest_1.0-4
[137] jmv_0.9.6.1	Rcpp_1.0.2
[139] car_3.0-3	broom_0.5.2
[141] metafor_2.1-0	ez_4.4-0
[143] BayesFactor_0.9.12-4.2	performance_0.3.0
[145] later_0.8.0	writexl_1.1
[147] httr_1.4.1	psych_1.8.12
[149] sjstats_0.17.6	colorspace_1.4-1
[151] rvest_0.3.4	splines_3.6.0
[153] expm_0.999-4	fit.models_0.5-14
[155] xtable_1.8-4	nloptr_1.2.1
[157] rstan_2.19.2	zeallot_0.1.0
[159] modeltools_0.2-22	R6_2.4.0
[161] broom.mixed_0.2.4	Hmisc_4.2-0
[163] pillar_1.4.2	htmltools_0.4.0
[165] mime_0.7	glue_1.3.1
[167] minqa_1.2.4	DT_0.9
[169] codetools_0.2-16	jomo_2.6-9
[171] pkgbuild_1.0.6	pcaPP_1.9-73
[173] mvtnorm_1.0-11	furrr_0.1.0

[175]	lattice_0.20-38	tibble_2.1.3
[177]	curl_4.2	DescTools_0.99.28
[179]	gtools_3.8.1	logspline_2.1.13
[181]	zip_2.0.4	openxlsx_4.1.0.1
[183]	survival_2.44-1.1	rmarkdown_1.16
[185]	munsell_0.5.0	rsample_0.0.5
[187]	sjmisc_2.8.1	haven_2.1.1
[189]	reshape2_1.4.3	gtable_0.3.0
[191]	bayestestR_0.3.0	

34 Notes

Last update on 2019-11-08 23:38:35

Serdar Balci, MD, Pathologist

drserdarbalci@gmail.com

<https://rpubs.com/sbalci/CV>