

Comparing means

Data Analysis Using R (2017)

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1 Two independent samples

1.1 Independent t-test

```
library(foreign)
library(psych)
cholest = read.spss("cholest.sav", to.data.frame = T)
str(cholest)

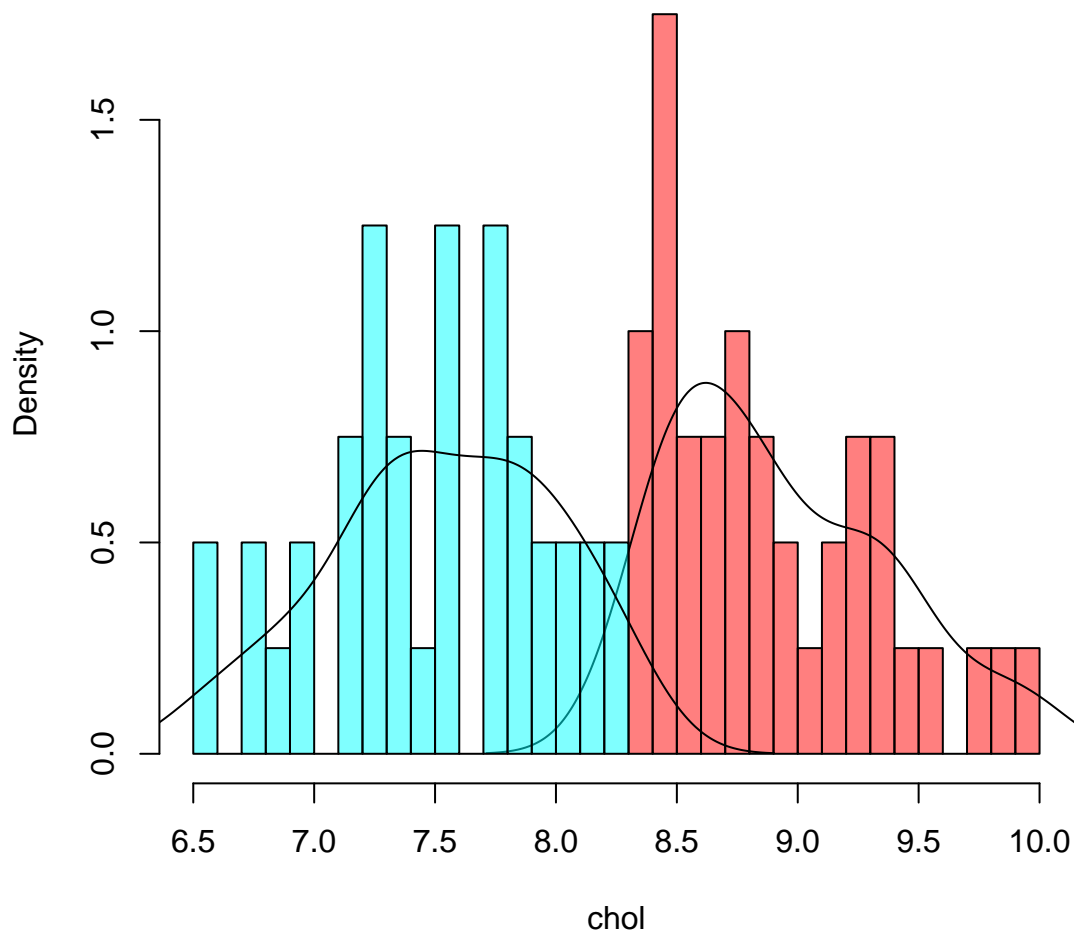
## 'data.frame': 80 obs. of 5 variables:
## $ chol : num 6.5 6.6 6.8 6.8 6.9 7 7 7.2 7.2 7.2 ...
## $ age : num 38 35 39 36 31 38 33 36 40 34 ...
## $ exercise: num 6 5 6 5 4 4 5 5 4 6 ...
## $ sex : Factor w/ 2 levels "female","male": 2 2 2 2 2 2 2 2 2 2 ...
## $ categ : Factor w/ 3 levels "Grp A","Grp B",...: 1 1 1 1 1 1 1 1 1 1 ...
## - attr(*, "variable.labels")= Named chr "cholesterol in mmol/L" "age in year" "duration of exercise" ...
## ..- attr(*, "names")= chr "chol" "age" "exercise" "sex" ...
## - attr(*, "codepage")= int 65001
```

```
head(cholest)
```

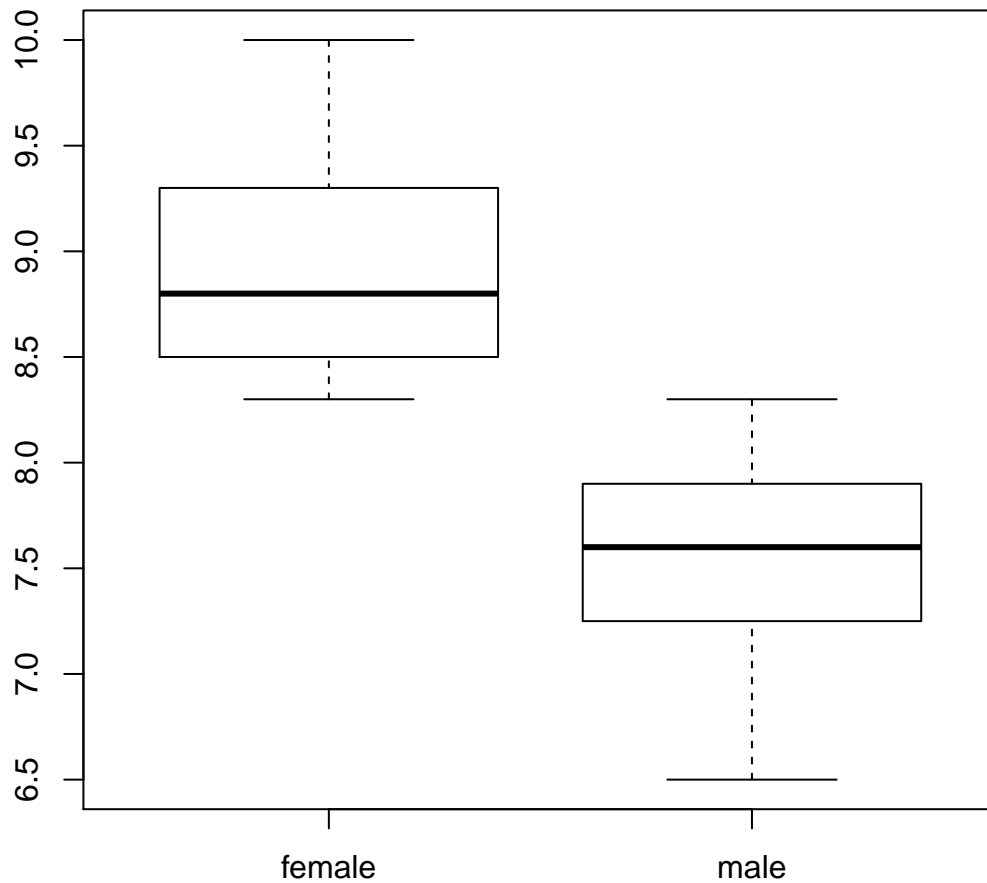
```
## chol age exercise sex categ
## 1 6.5 38      6 male Grp A
## 2 6.6 35      5 male Grp A
## 3 6.8 39      6 male Grp A
## 4 6.8 36      5 male Grp A
## 5 6.9 31      4 male Grp A
## 6 7.0 38      4 male Grp A
```

```
histBy(cholest, "chol", group = "sex")
```

Histograms by group



```
boxplot(chol ~ sex, data = cholest)
```



```
t.test(chol ~ sex, data = cholest)
```

```
##
## Welch Two Sample t-test
##
## data: chol by sex
## t = 13.504, df = 77.933, p-value < 2.2e-16
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
##  1.189337 1.600663
## sample estimates:
## mean in group female   mean in group male
##                8.9275                7.5325
```

```
# ?t.test # other options
```

1.2 Mann-Whitney U test / Wilcoxon rank-sum test

```
wilcox.test(chol ~ sex, data = cholest) # not accurate for ties

## Warning in wilcox.test.default(x = c(8.3, 8.3, 8.4, 8.4, 8.5, 8.5, 8.5, : cannot compute
## exact p-value with ties

##
## Wilcoxon rank sum test with continuity correction
##
## data: chol by sex
## W = 1598, p-value = 1.568e-14
## alternative hypothesis: true location shift is not equal to 0
# ?wilcox.test
library(coin)

## Loading required package: survival
wilcox_test(chol ~ sex, data = cholest)

##
## Asymptotic Wilcoxon-Mann-Whitney Test
##
## data: chol by sex (female, male)
## Z = 7.6867, p-value = 1.51e-14
## alternative hypothesis: true mu is not equal to 0
wilcox_test(chol ~ sex, data = cholest, distribution = "exact")

##
## Exact Wilcoxon-Mann-Whitney Test
##
## data: chol by sex (female, male)
## Z = 7.6867, p-value < 2.2e-16
## alternative hypothesis: true mu is not equal to 0
# ?wilcox_test
```

2 Two dependent samples

2.1 Paired t-test

```
sbp = read.spss("sbp.sav", to.data.frame = T)
t.test(sbp$S1, sbp$S2, paired = T)

##
## Paired t-test
##
## data: sbp$S1 and sbp$S2
## t = -0.81954, df = 10, p-value = 0.4316
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -5.071058 2.343785
## sample estimates:
```

```
## mean of the differences
##          -1.363636
```

2.2 Wilcoxon signed-rank test

```
wilcox.test(sbp$S1, sbp$S2, paired = T)
```

```
## Warning in wilcox.test.default(sbp$S1, sbp$S2, paired = T): cannot compute exact p-value
## with ties
```

```
## Warning in wilcox.test.default(sbp$S1, sbp$S2, paired = T): cannot compute exact p-value
## with zeroes
```

```
##
## Wilcoxon signed rank test with continuity correction
##
## data: sbp$S1 and sbp$S2
## V = 3, p-value = 0.5708
## alternative hypothesis: true location shift is not equal to 0
```

```
wilcoxsign_test(sbp$S1 ~ sbp$S2)
```

```
##
## Asymptotic Wilcoxon-Pratt Signed-Rank Test
##
## data: y by x (pos, neg)
## stratified by block
## Z = -0.94346, p-value = 0.3454
## alternative hypothesis: true mu is not equal to 0
```

```
wilcoxsign_test(sbp$S1 ~ sbp$S2, distribution = "exact")
```

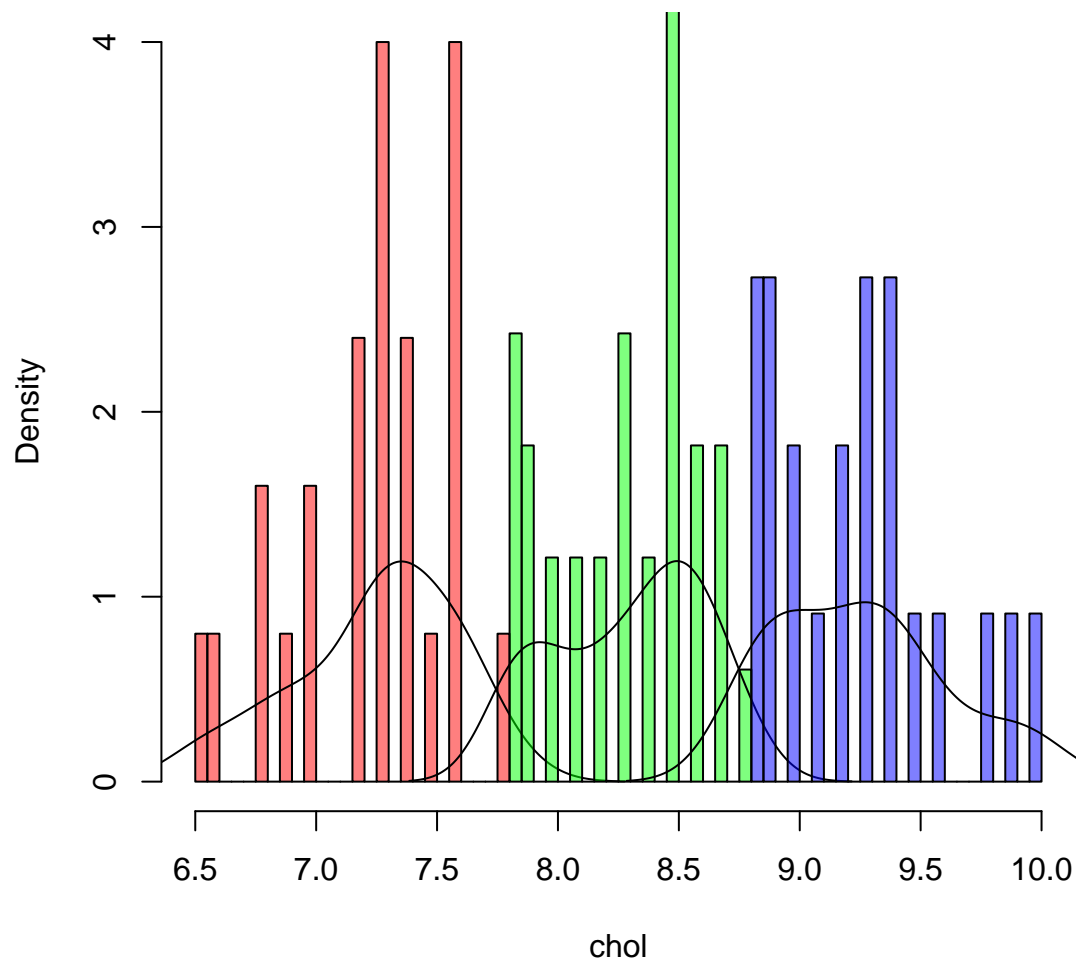
```
##
## Exact Wilcoxon-Pratt Signed-Rank Test
##
## data: y by x (pos, neg)
## stratified by block
## Z = -0.94346, p-value = 0.625
## alternative hypothesis: true mu is not equal to 0
```

3 Independent samples

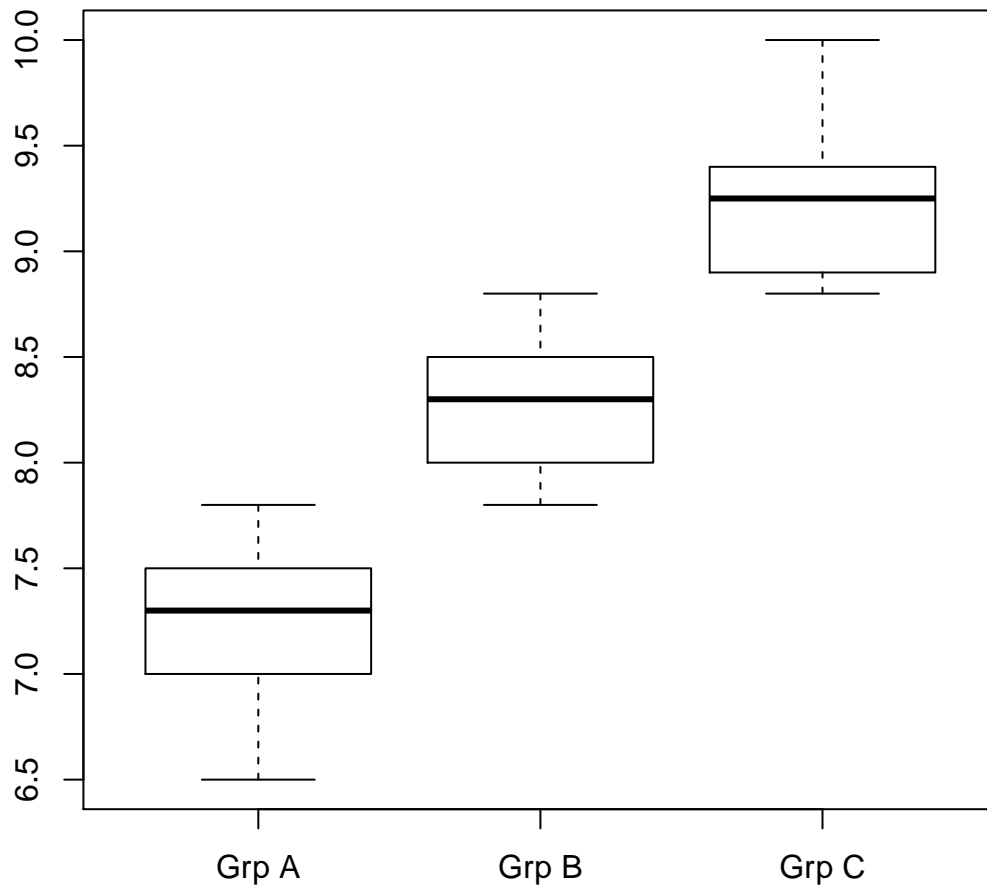
3.1 One-way ANOVA

```
histBy(cholest, "chol", group = "categ")
```

Histograms by group



```
boxplot(chol ~ categ, data = cholest)
```



```
aov_chol = aov(chol ~ categ, data = cholest)
summary(aov_chol)
```

```
##           Df Sum Sq Mean Sq F value Pr(>F)
## categ      2  47.13   23.57  215.1 <2e-16 ***
## Residuals  77   8.44    0.11
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

3.2 Kruskal-Wallis test

```
kruskal.test(chol ~ categ, data = cholest)
```

```
##
## Kruskal-Wallis rank sum test
##
```

```
## data: chol by categ
## Kruskal-Wallis chi-squared = 69.188, df = 2, p-value = 9.464e-16
```

4 Dependent samples

4.1 Repeated measures ANOVA

```
library(car)

##
## Attaching package: 'car'
## The following object is masked from 'package:psych':
##
##   logit

time = ordered(rep(1:3))
idesign = data.frame(time)
model_rm = lm(cbind(S1, S2, S3) ~ 1, sbp)
aov_rm = Anova(model_rm, idata = idesign, idesign = ~time)

## Note: model has only an intercept; equivalent type-III tests substituted.
summary(aov_rm, multivariate = F) # univariate approach

## Warning in summary.Anova.mlm(aov_rm, multivariate = F): HF eps > 1 treated as 1

##
## Univariate Type III Repeated-Measures ANOVA Assuming Sphericity
##
##           SS num Df Error SS den Df      F    Pr(>F)
## (Intercept) 487276      1  5824.2    10 836.6337 5.686e-11 ***
## time          29      2   271.2    20  1.0615  0.3647
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
##
## Mauchly Tests for Sphericity
##
##      Test statistic p-value
## time      0.91149 0.65899
##
##
## Greenhouse-Geisser and Huynh-Feldt Corrections
## for Departure from Sphericity
##
##      GG eps Pr(>F[GG])
## time 0.91869  0.3608
##
##      HF eps Pr(>F[HF])
## time 1.115516 0.3646518

summary(aov_rm) # multivariate approach

## Warning in summary.Anova.mlm(aov_rm): HF eps > 1 treated as 1
```



```

##
## Type III Repeated Measures MANOVA Tests:
##
## -----
##
## Term: (Intercept)
##
## Response transformation matrix:
## (Intercept)
## S1 1
## S2 1
## S3 1
##
## Sum of squares and products for the hypothesis:
## (Intercept)
## (Intercept) 1461827
##
## Multivariate Tests: (Intercept)
## Df test stat approx F num Df den Df Pr(>F)
## Pillai 1 0.98819 836.6337 1 10 5.6855e-11 ***
## Wilks 1 0.01181 836.6337 1 10 5.6855e-11 ***
## Hotelling-Lawley 1 83.66337 836.6337 1 10 5.6855e-11 ***
## Roy 1 83.66337 836.6337 1 10 5.6855e-11 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## -----
##
## Term: time
##
## Response transformation matrix:
## time.L time.Q
## S1 -7.071068e-01 0.4082483
## S2 -7.850462e-17 -0.8164966
## S3 7.071068e-01 0.4082483
##
## Sum of squares and products for the hypothesis:
## time.L time.Q
## time.L 4.545455 10.49728
## time.Q 10.497278 24.24242
##
## Multivariate Tests: time
## Df test stat approx F num Df den Df Pr(>F)
## Pillai 1 0.153110 0.8135593 2 9 0.47339
## Wilks 1 0.846890 0.8135593 2 9 0.47339
## Hotelling-Lawley 1 0.180791 0.8135593 2 9 0.47339
## Roy 1 0.180791 0.8135593 2 9 0.47339
##
## Univariate Type III Repeated-Measures ANOVA Assuming Sphericity
##
## SS num Df Error SS den Df F Pr(>F)
## (Intercept) 487276 1 5824.2 10 836.6337 5.686e-11 ***
## time 29 2 271.2 20 1.0615 0.3647
## ---

```

```

## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
##
## Mauchly Tests for Sphericity
##
##      Test statistic p-value
## time      0.91149 0.65899
##
##
## Greenhouse-Geisser and Huynh-Feldt Corrections
## for Departure from Sphericity
##
##      GG eps Pr(>F[GG])
## time 0.91869      0.3608
##
##      HF eps Pr(>F[HF])
## time 1.115516   0.3646518

```

4.2 Friedman test

```
friedman.test(as.matrix(sbp[, c("S1", "S2", "S3")]))
```

```

##
## Friedman rank sum test
##
## data:  as.matrix(sbp[, c("S1", "S2", "S3")])
## Friedman chi-squared = 1.2381, df = 2, p-value = 0.5385

```

References

- Fox, J., & Weisberg, S. (2017). *Car: Companion to applied regression*. Retrieved from <https://CRAN.R-project.org/package=car>
- Hothorn, T., Hornik, K., van de Wiel, M. A., Winell, H., & Zeileis, A. (2017). *Coin: Conditional inference procedures in a permutation test framework*. Retrieved from <https://CRAN.R-project.org/package=coin>
- Revelle, W. (2017). *Psych: Procedures for psychological, psychometric, and personality research*. Retrieved from <https://CRAN.R-project.org/package=psych>