

Introduction to R

Holger Sennhenn-Reulen^a, Nordwestdeutsche Forstliche Versuchsanstalt (NW-FVA)

July 07, 2021 (Version 0.1)

Contents

1	Introduction: What is R?	2
2	Data	2
	Drought	2
3	Section 1	4
	Outcomes	4
	3.1 Tree species	4

All contents are licensed under CC BY-NC-ND 4.0 ([Link](#)).

^aPrivate webpage: uncertaintree.github.io

1 Introduction: What is R?

...

2 Data

We will use two data-sets within this course.

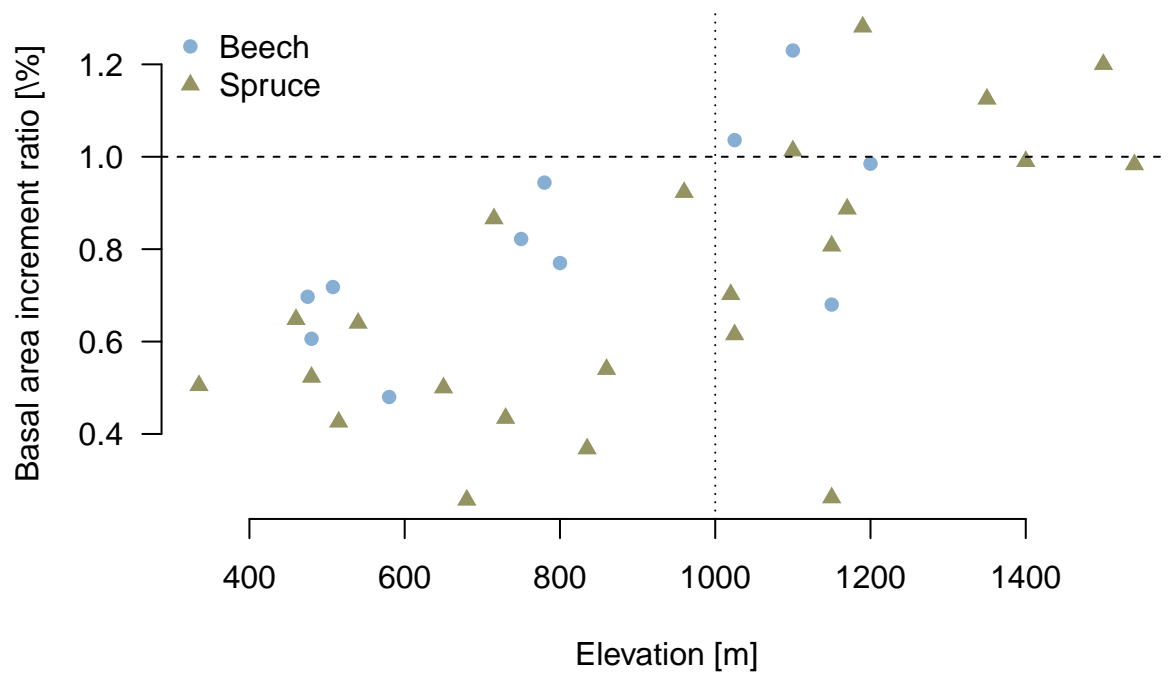
Drought

```
bair <- c(.505, .648, .523, .426, .64, .5, .257, .866, .434, .368, .54, .923, .702,
        .615, 1.013, .807, .262, .887, 1.281, 1.125, .99, 1.2, .983, .697, .606,
        .718, .48, .822, .944, .77, 1.036, 1.23, .68, .985)
elev <- c(335, 460, 480, 515, 540, 650, 680, 715, 730, 835, 860, 960,
        1020, 1025, 1100, 1150, 1150, 1170, 1190, 1350, 1400, 1500, 1540,
        475, 480, 507.5, 580, 750, 780, 800, 1025, 1100, 1150, 1200)
species <- c("Spruce", "Spruce", "Spruce", "Spruce", "Spruce", "Spruce",
            "Spruce", "Spruce", "Spruce", "Spruce", "Spruce", "Spruce", "Spruce",
            "Spruce", "Spruce", "Spruce", "Spruce", "Spruce", "Spruce", "Spruce",
            "Spruce", "Spruce", "Spruce", "Beech", "Beech", "Beech", "Beech",
            "Beech", "Beech", "Beech", "Beech", "Beech", "Beech", "Beech")
drought <- data.frame(bair = bair,
                     elev = elev,
                     species = species)
summary(drought)
```

##	bair	elev	species
## Min.	:0.2570	Min. : 335.0	Beech :11
## 1st Qu.	:0.5272	1st Qu.: 597.5	Spruce:23
## Median	:0.7100	Median : 847.5	
## Mean	:0.7489	Mean : 888.3	
## 3rd Qu.	:0.9732	3rd Qu.:1150.0	
## Max.	:1.2810	Max. : 1540.0	

Plot

```
pch_here <- c(16, 17)
col_here <- colorspace::diverging_hcl(n = 20, pal = "Lisbon")[c(4, 16)]
plot(drought$elev, drought$bair, las = 1, bty = "n",
     xlab = "Elevation [m]", ylab = "Basal area increment ratio [\\%]",
     pch = pch_here[1 + (drought$species == "Spruce")],
     col = col_here[1 + (drought$species == "Spruce")])
abline(h = 1, lty = 2)
abline(v = 1000, lty = 3)
legend("topleft", pch = pch_here, col = col_here, legend = c("Beech", "Spruce"),
     bg = "white", bty = "n")
```



A Simple Model

```
m <- lm(bair ~ elev*species, data = drought)
summary(m)

##
## Call:
## lm(formula = bair ~ elev * species, data = drought)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -0.58062 -0.07711  0.01943  0.14246  0.41573
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)   3.969e-01  2.035e-01   1.951  0.0605 .
## elev          5.201e-04  2.403e-04   2.164  0.0386 *
## speciesSpruce -2.056e-01  2.397e-01  -0.858  0.3979
## elev:speciesSpruce 4.619e-05  2.723e-04   0.170  0.8664
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.2106 on 30 degrees of freedom
## Multiple R-squared:  0.463, Adjusted R-squared:  0.4093
## F-statistic: 8.621 on 3 and 30 DF, p-value: 0.0002811
```

3 Section 1

Outcomes

Definition (Outcome): The *outcome* of a random variable X is a value that the random variable might (in theory/model) take on when applying a measurement method.

3.1 Tree species

3.1.0.1 Example (Tree species)

Random selection of a single tree from the last German National Forest Inventory data-base builds a random variable X – for characteristic *tree species group*ⁱ – with sample space Ω :

$$\Omega = \{\text{oak, beech, odtwle, odtwsle, spruce, fir, douglas fir, pine, larch}\}.$$

All possible ω are:

$$\omega = \{\text{oak}\}, \omega = \{\text{beech}\}, \dots, \omega = \{\text{larch}\}.$$

Random variables are actually only truly helpful if we can define them such that mathematical manipulations can be applied on them as we do with real numbers. So instead of X , we better define eight indicators [Kadane, 2020] X_1, \dots, X_8 with samples spaces $\Omega_1 = \{\text{not oak, oak}\}, \Omega_2 = \{\text{not beech, beech}\}, \dots, \Omega_8 = \{\text{not larch, larch}\}$:

$$X_1 = \begin{cases} 0, & \omega = \{\text{not oak}\} \\ 1, & \omega = \{\text{oak}\} \end{cases}, X_2 = \begin{cases} 0, & \omega = \{\text{not beech}\} \\ 1, & \omega = \{\text{beech}\} \end{cases}, \dots, X_8 = \begin{cases} 0, & \omega = \{\text{not larch}\} \\ 1, & \omega = \{\text{larch}\} \end{cases},$$

and thus with outcomes $x_j = 0$, or $x_j = 1$, $j = 1, \dots, 8$.

For repeated random experiments, such outcomes are stored in the form of a vector – columns of a data-set in R.

3.1.0.2 Example (Tree species) continued

Random selection of $n = 10$ trees results in a vector:

$$\mathbf{x}_1 = (0, 0, 1, 0, 0, 0, 0, 0, 0, 0)^\top$$

with $x_{1,1} = 0$, $x_{1,2} = 0$, $x_{1,3} = 1$, \dots , $x_{1,10} = 0$.

3.1.0.3 Example (Drought)

The *drought* data contains measures on the ratio of this year's basal area increment (BAI) in relation to last year's BAI: % So BAI ratios X :

$$X = \frac{\text{BAI this year}}{\text{BAI last year}}$$

BAI measures can be positive and negativeⁱⁱ, but in order to avoid $-1 = \frac{-1}{1} = \frac{1}{-1}$, we take a 'growing process definition'ⁱⁱⁱ by which basal area increments become strictly positive. Consequently, the sample space of X is the positive real line, $\Omega = \mathbb{R}_+$, with samples such as:

$$\omega = \{1.37\}, \omega = \{0.93\}, \dots$$

and positive real valued outcomes such as:

$$\mathbf{x} = (1.37, .93, \dots)^\top$$

ⁱodtwle: other deciduous trees with long life expectancy, odtwsle: other deciduous trees with short life expectancy

ⁱⁱ<https://www.sciencedirect.com/science/article/pii/S0378112720314948#t0005>

ⁱⁱⁱ<http://www.personal.psu.edu/agl/Johnson&Abrams%20TRACE.pdf>

References

J.B. Kadane. *Principles of Uncertainty*. Chapman and Hall/CRC, 2020. ISBN 9781138052734.